

Fluctuation Spectroscopy

Making signal out of noise

Don C. Lamb

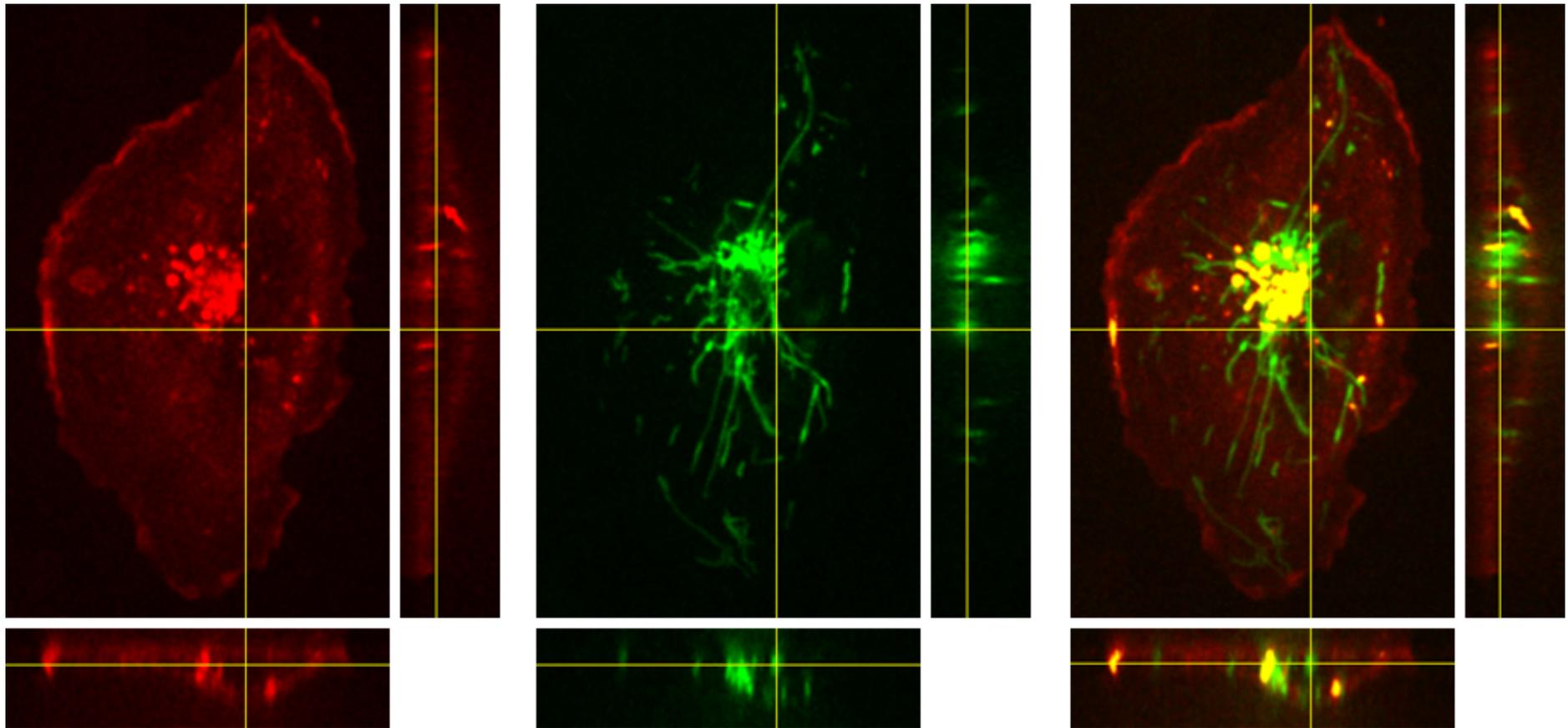


Laboratory for Fluorescence
Applications in **B**iological Systems
Institute of Physical Chemistry
Munich, Germany



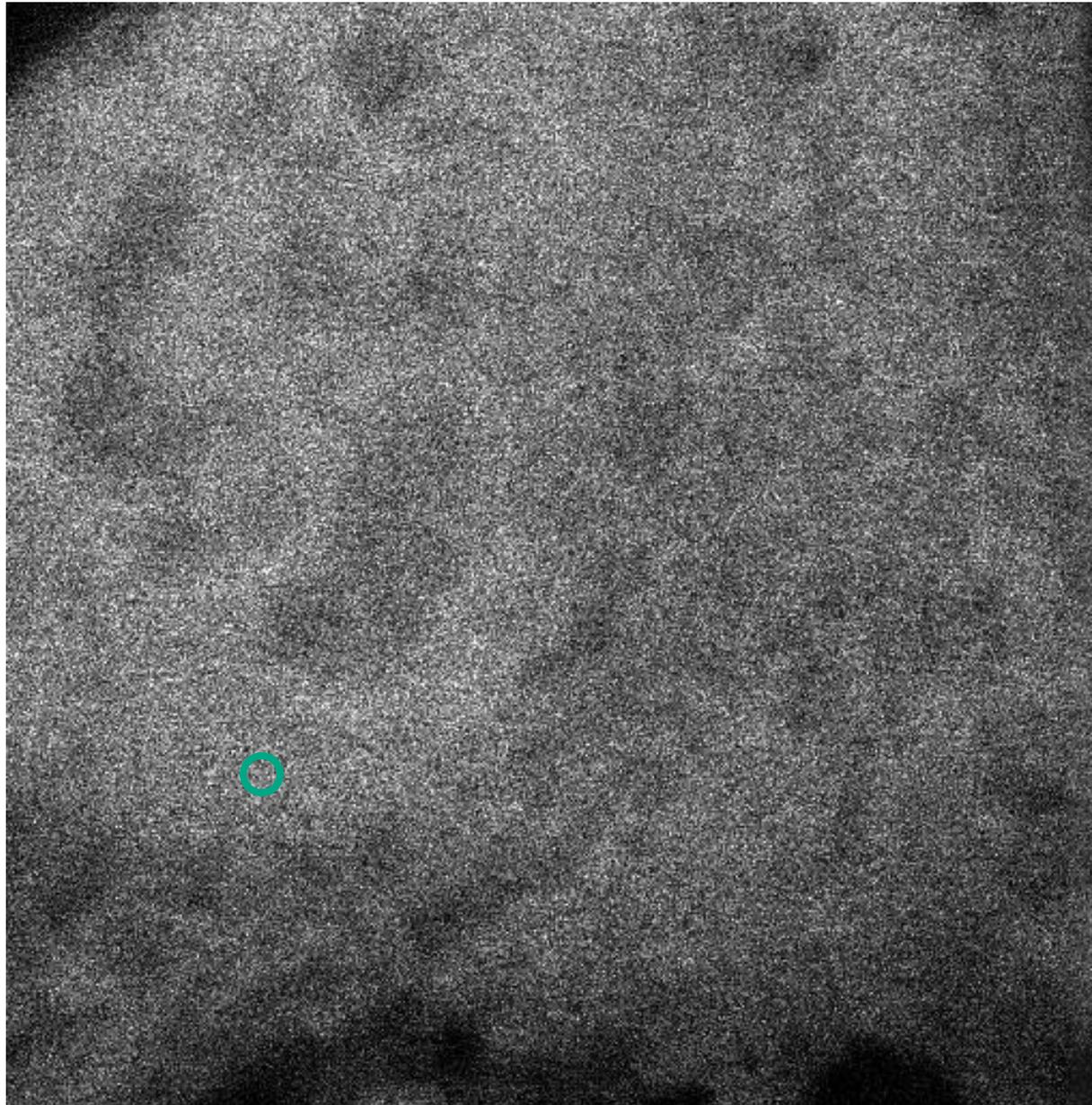


Microscopy: What we show you



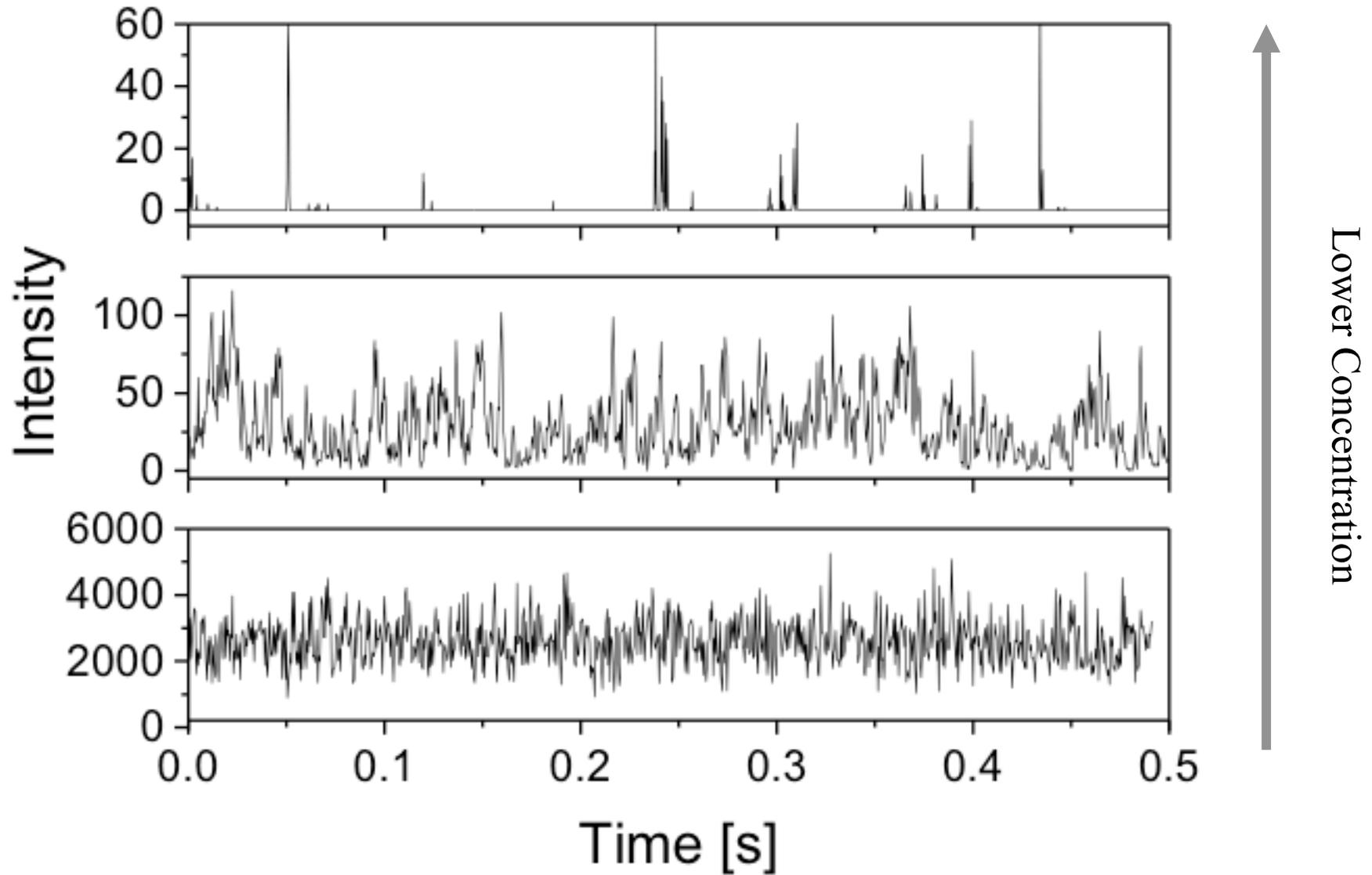


Microscopy: What we measure in living cells



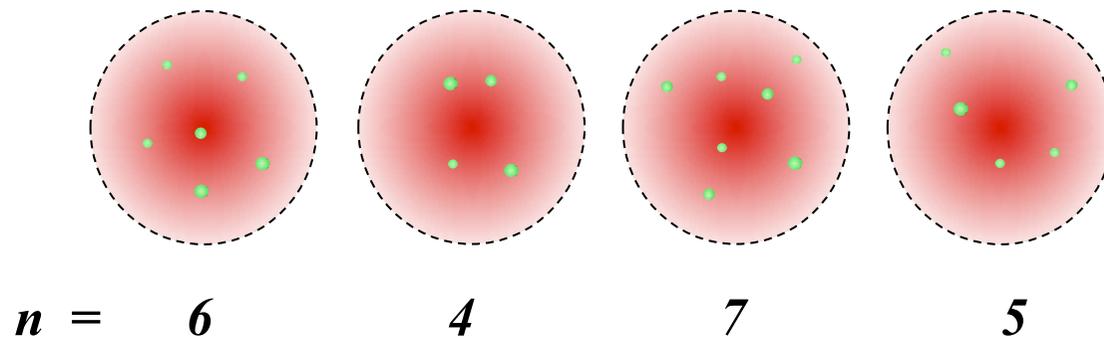


Fluctuation Measurements





Fluctuations in molecular number in a small volume.



$$P(N) = \frac{\langle N \rangle^N e^{-\langle N \rangle}}{N!}$$

$$\langle \Delta N^2 \rangle = \langle N \rangle$$

A Poissonian Process

Experiments that result in counting the number of events in a given time or in a given object can be described by a Poisson process provided:

- Number changes on nonoverlapping intervals are independent.
- The probability of exactly one change occurring in a sufficiently short interval of length h is approximately λh
- The probability of two or more changes in a sufficiently short interval is essentially zero.

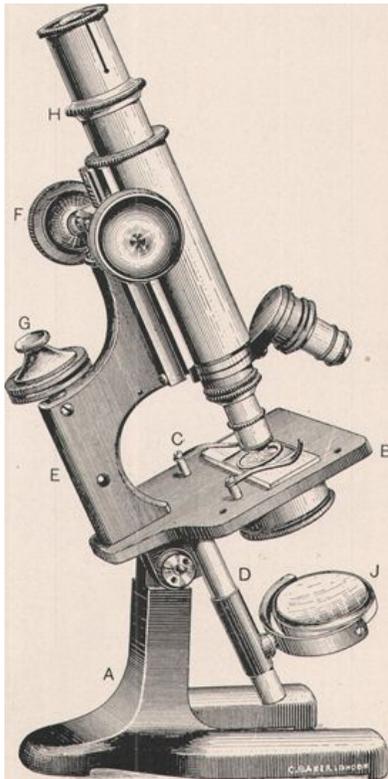


Fluctuations



Observation of gold colloids using an ultra-microscope (Svedberg and Inouye, *Zeitschr f. Physik Chemie* **1911**, 77:145-119)

Measurement of the Equilibrium Thermodynamic Fluctuations in Molecular Number

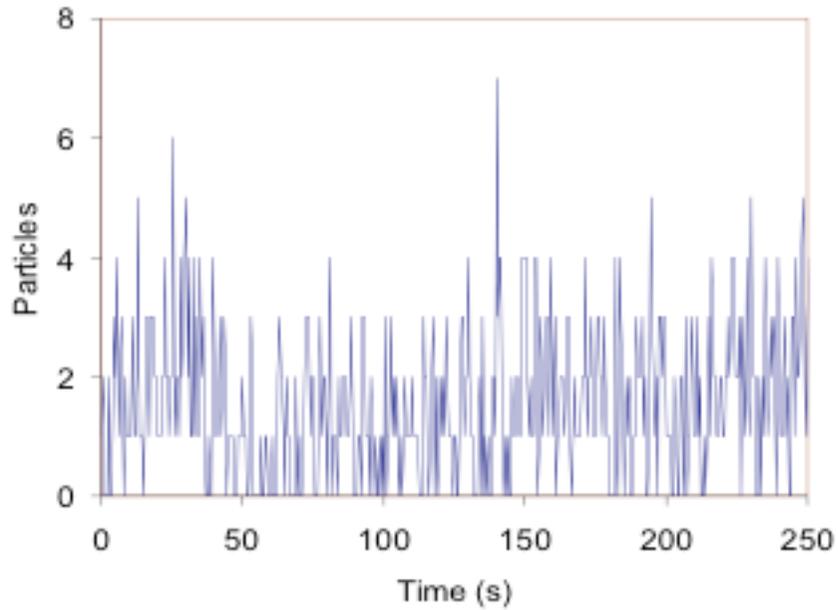


1200020013241231021111311251110233133322111224221226122
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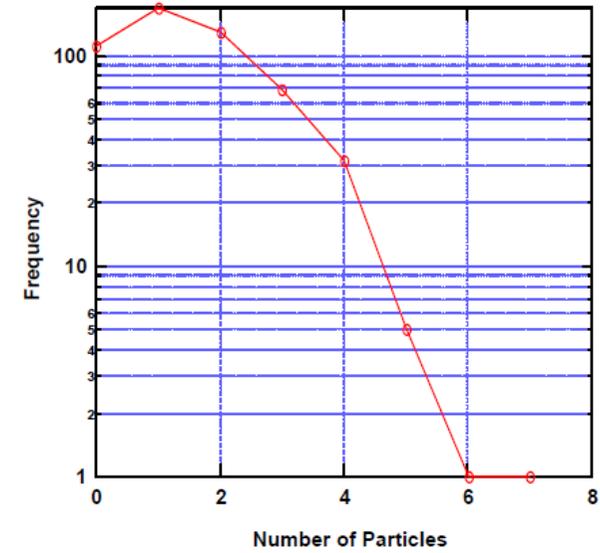
<http://www.1911encyclopedia.org/Microscope>



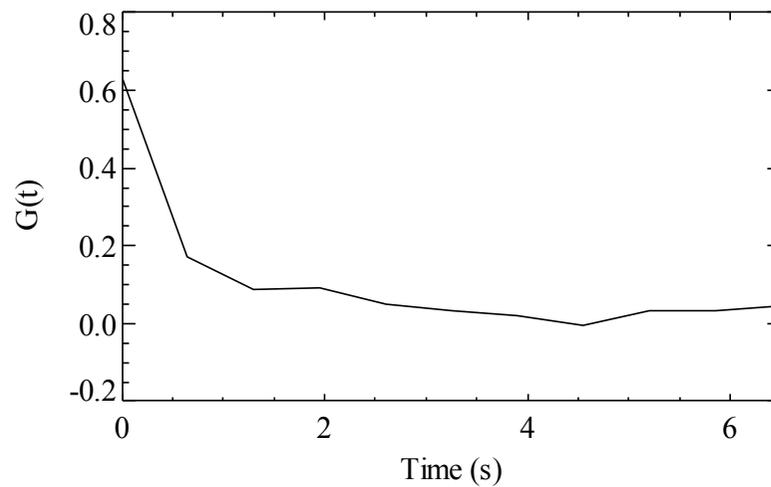
Autocorrelation Analysis



PCH



Autocorrelation Analysis



$$\langle N \rangle = 1.55 \text{ particles}$$

$$\tau \sim 1.5 \text{ s}$$



Autocorrelation Analysis



The normalized autocorrelation function (ACF) is given by:

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

where

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

For processes that are:

Stationary: i.e. the average parameters do not change with time

the ACF is independent of the absolute time

Ergodic: i.e. every sizeable sampling of the process is representative of the whole

the time average is equal to the ensemble average

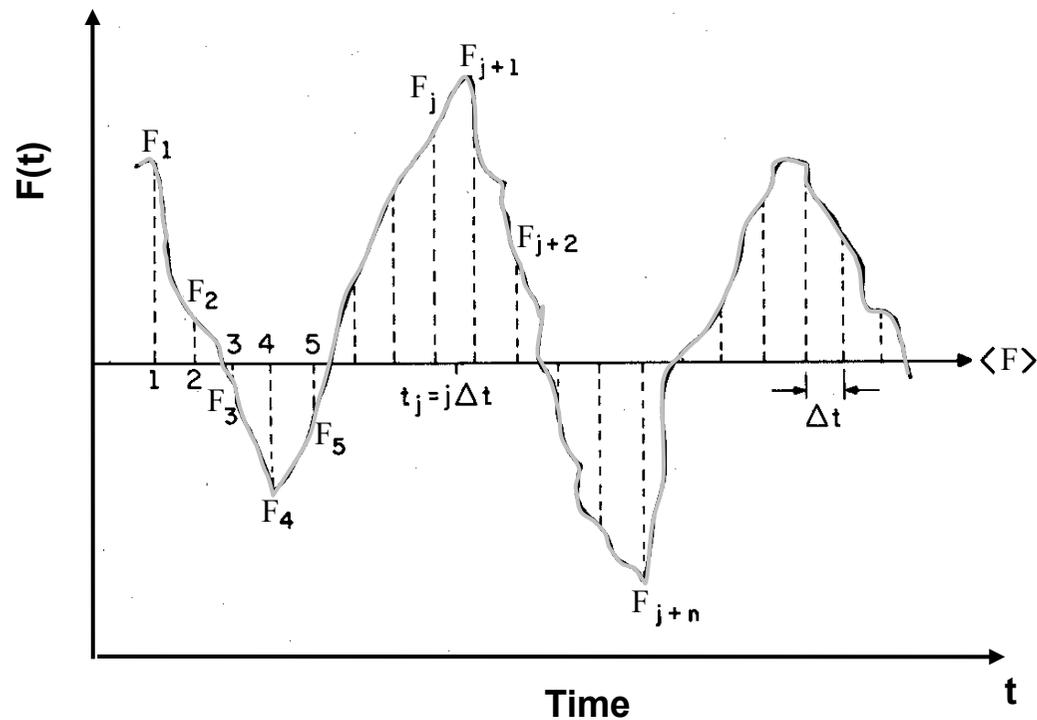
$$\frac{\langle \delta F(t)\delta F(t+\tau) \rangle}{\langle F(t) \rangle^2} = \frac{\langle \delta F(0)\delta F(\tau) \rangle}{\langle F \rangle^2}$$



Properties of the Autocorrelation Function



The autocorrelation function (ACF) measures the self similarity of the observable F as a function of t



$$G(\tau) = \frac{\langle \delta F(0) \delta F(\tau) \rangle}{\langle F \rangle^2}$$
$$= \frac{\sum_{i=1}^{\ell} \delta F_i \delta F_{i+\tau} / \ell}{\left(\sum_{i=1}^{\ell} F_i / \ell \right)^2}$$

$$(\delta F_i)^2 \geq 0$$

$$\delta F_i \delta F_{i+\tau} \text{ can be } < 0$$

$G(\tau)$ has maximum at $G(0)$



Properties of the Autocorrelation Function



The amplitude is proportional to the size of the fluctuations

$$G(0) = \frac{\langle \delta F(t) \delta F(t) \rangle}{\langle F \rangle^2} = \frac{\langle \delta F(t)^2 \rangle}{\langle F \rangle^2} = \frac{\langle (F(t) - \langle F(t) \rangle)^2 \rangle}{\langle F \rangle^2} = \frac{\sum_{i=1}^{\ell} (F_i - \langle F \rangle)^2 / \ell}{\left(\sum_{i=1}^{\ell} F_i / \ell \right)^2}$$

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

For non-conserved, non-periodic signals

$G(t) \rightarrow 0$ as $t \rightarrow \infty$

For a Poissonian process

Alternately, the ACF is sometimes given as:

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

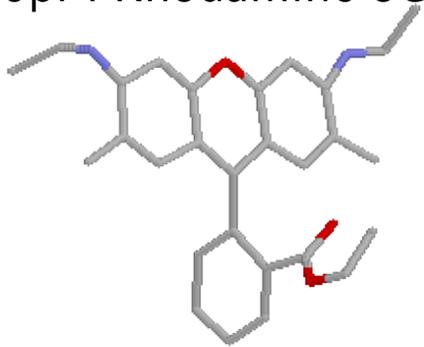
$g(\tau)$ can be interpreted as being proportional to the probability of detecting a photon at delay τ when a photon was detected at $\tau = 0$



ACF of Freely Diffusing Molecules



230pM Rhodamine 6G in buffer



Amplitude ACF \Rightarrow

Concentration

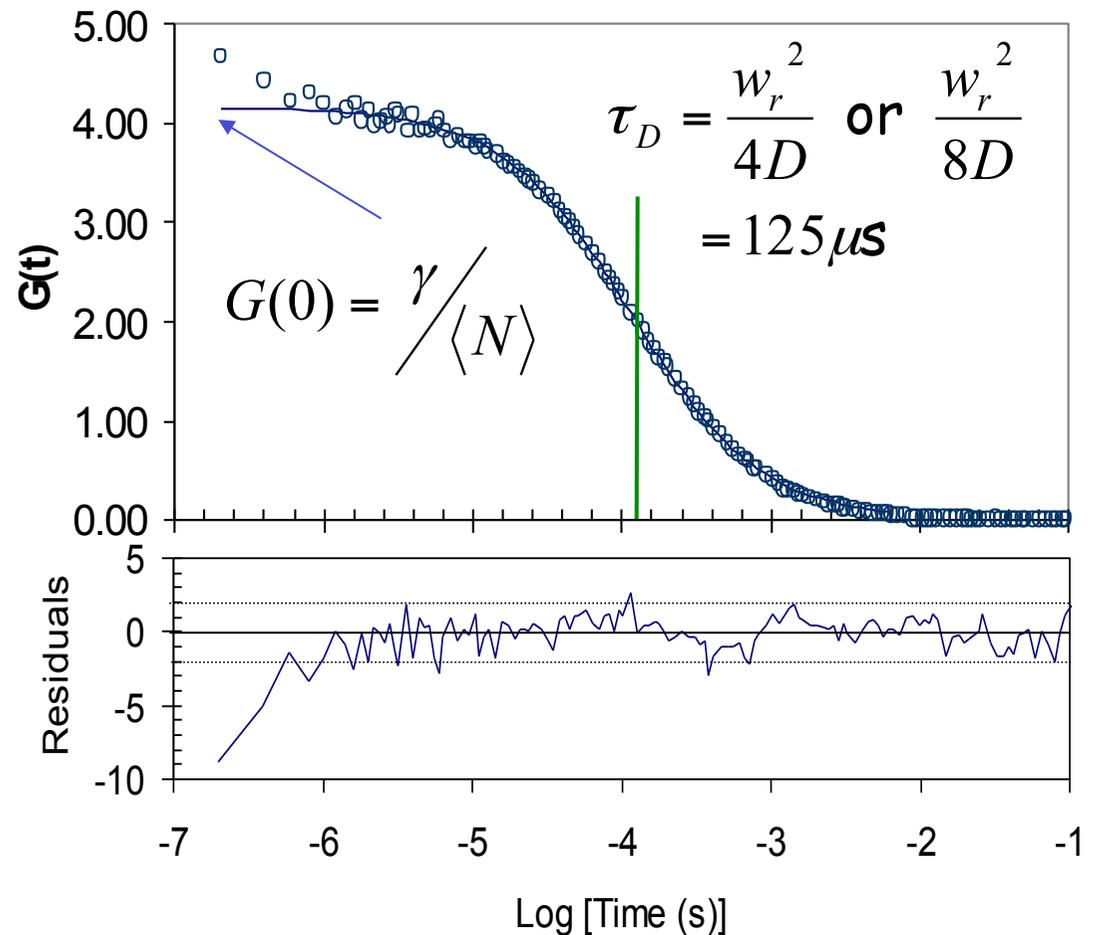
$\langle N \rangle = 0.085$ molecules

Decay ACF \Rightarrow

Diffusion Constant

$D = 415 \mu\text{m}^2/\text{s}$ ($w_r = 456 \text{ nm}$)

$$G_D(\tau, N, \tau_D) = \frac{\gamma}{\langle N \rangle} \left(\frac{1}{1 + \tau/\tau_D} \right) \left(\frac{1}{1 + (w_r/w_z)^2 \tau/\tau_D} \right)^{1/2}$$

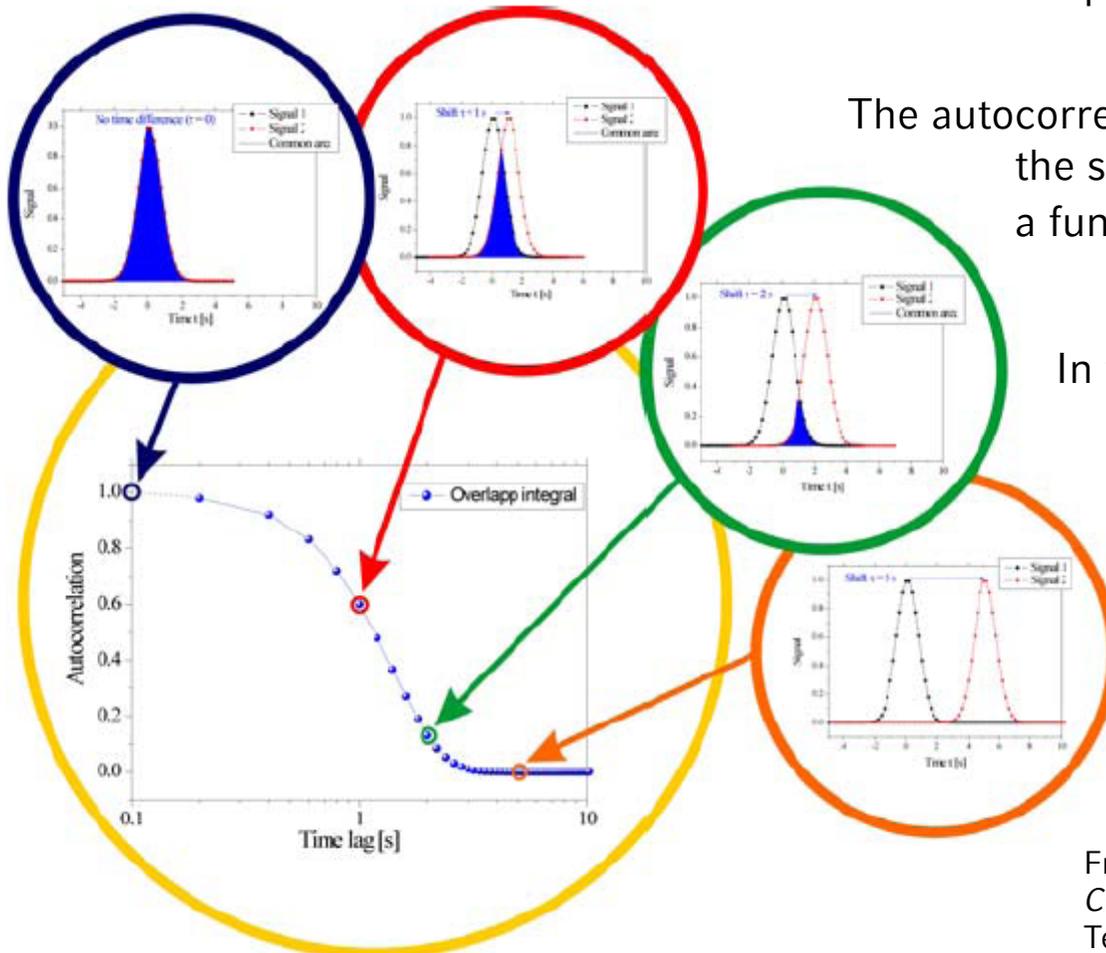


$$G(\tau) = \frac{\langle F(t)F(t+\tau) \rangle - \langle F(t) \rangle^2}{\langle F(t) \rangle^2}$$

Photons are not detected stochastically, but in bursts when a molecule transverses the probe volume

The autocorrelation function (ACF) measures the self similarity of the time trace as a function of the shift τ

In this case, the ACF measures the averaged duration of a burst of photons



Why Spectroscopy?

Spectroscopy of the Fluctuations

From: Schwille, Haustein, *Fluorescence Correlation Spectroscopy*, In: Single Molecule Techniques, Biophysics Textbook Online



Signal-to-Noise Considerations



High Intensity Limit:

Uncertainty dominated by number of fluctuations:

$$\frac{S}{N} \approx \left(\frac{t_{\text{exp}}}{\tau_C} \right)^{1/2}$$

where t_{exp} is the measurement time of the experiment and τ_C is the correlation time of the fluctuations

Low Intensity Limit:

Uncertainty dominated by number of photons:

$$\frac{S}{N} \approx (t_{\text{exp}})^{1/2} I_T \frac{\gamma}{\langle N \rangle}$$

$$I_T = \varepsilon \langle N \rangle$$

$$\frac{S}{N} \approx (t_{\text{exp}})^{1/2} \varepsilon \gamma$$

Only possibilities to improve the S/N ratio are:

- extend the measurement time
- increase the counts per molecule second
- change the geometry

S/N is independent of sample concentration!!!



Limitations



Time Scale: ns/ μ s \rightarrow ms/s/hrs

Early time limit:

Detector afterpulsing: (100 ns - 5 μ s)

Detector deadtime: (2 ns - 30 ns)

Numbers of available photons:
(10 ns - 100 ns)

Long time limit:

Time molecule remains in
the excitation volume (typically \sim 1 ms)

Increase the long time limit by:

Increasing the excitation volume: (10 ms)

Placing sample in viscous solvents or gels: (s)

Slow reactions can be measured by changes in the
ACF with time. (hrs)

Concentration Limits:

\sim 200nM \rightarrow 1pM

Maximum Concentration: (200nM)

Detector Saturation

Other noise sources become
comparable to the signal

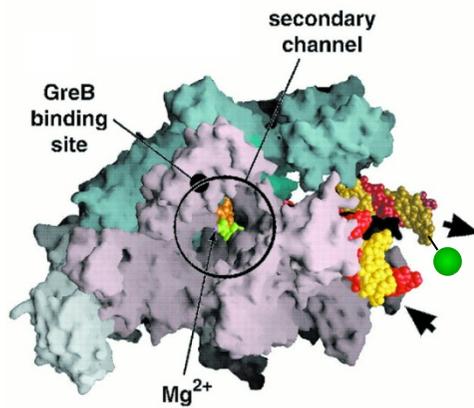
Minimum Concentration: (1pM)

Limit statistics

Impurities

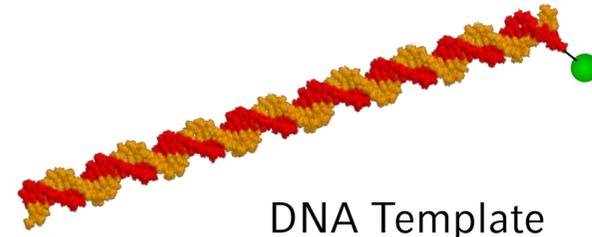


Molecular Interactions

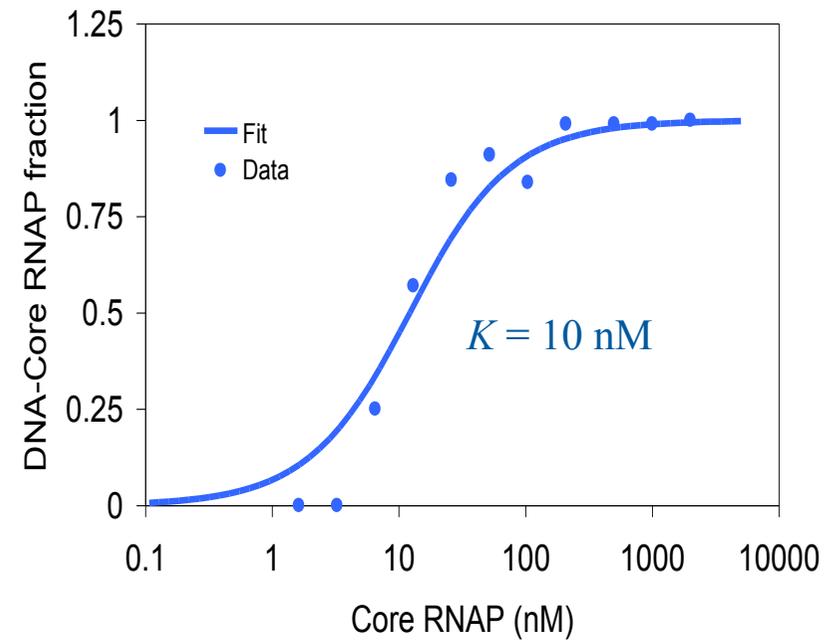
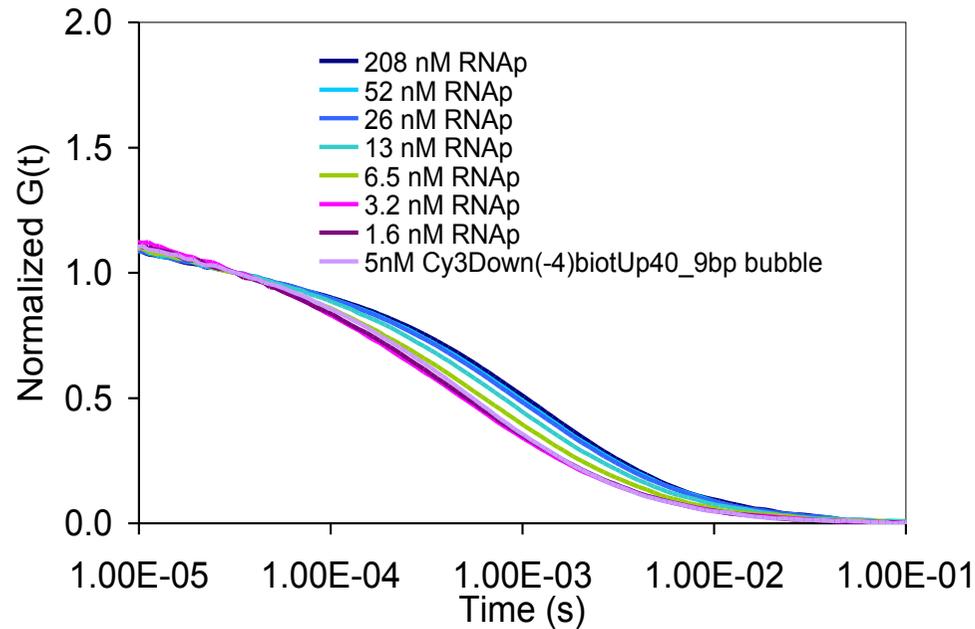


Titration and slow reactions resulting in a change in the diffusion constant can be followed using FCS

RNA Polymerase

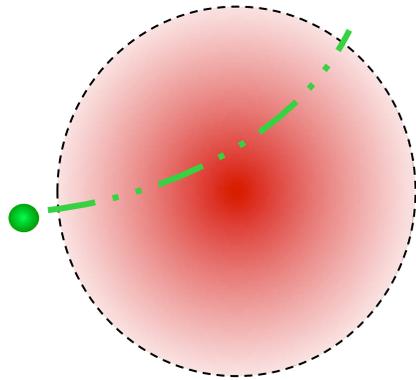


Titration of RNA Polymerase and DNA w/ 9 bp Artificial Bubble





Triplet State Blinking



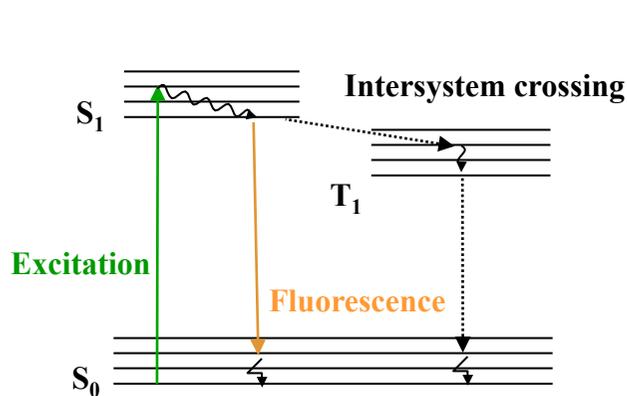
Freely diffusing, non-interacting particles in an open volume.

Photons are not detected stochastically, but in bursts when a molecule transverses the probe volume.

Several processes other than diffusion can lead to fluctuations in fluorescence intensity

e.g. Excitation into the Triplet State

If a particle blinks as it diffuses across the probe volume, an additional term appears in the fluctuation amplitude.



$$G(\tau) = \frac{\gamma}{\langle N \rangle} \left(\frac{1}{1 + \tau / \tau_D} \right) \left(\frac{1}{1 + (w_r / w_z)^2 \tau / \tau_D} \right)^{1/2} \left(1 + \frac{T}{1 - T} e^{-\frac{t}{\tau_T}} \right)$$

T is the fraction of molecules (on average) in the triplet state

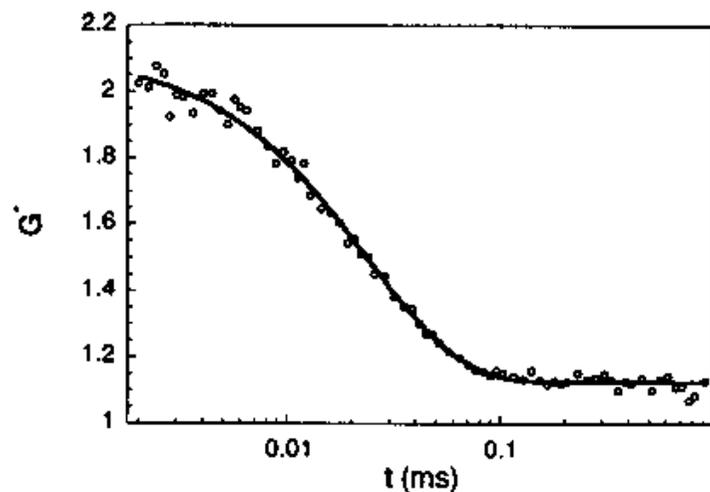
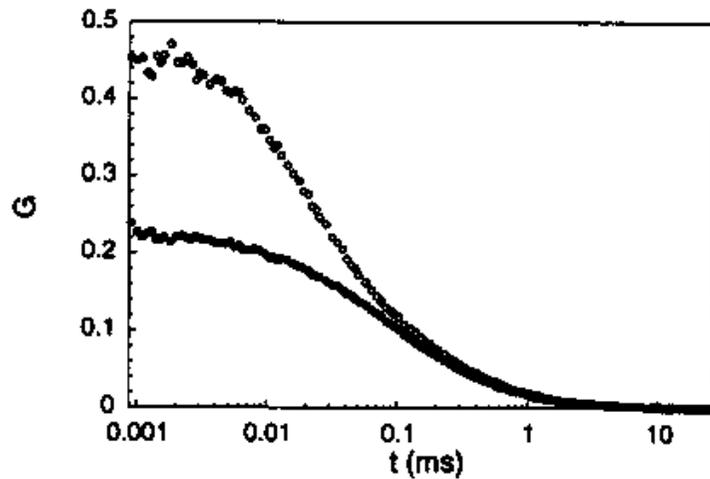
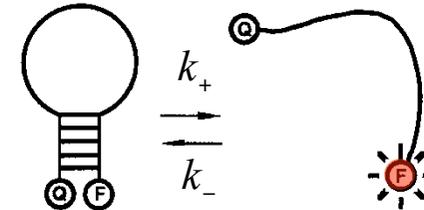
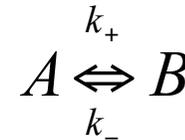
τ_T is the triplet lifetime.



Unimolecular Reaction



Dynamics of DNA Hairpin Formation



$$G(\tau) = G_D(\tau, N_A + N_B, \tau_D) \left[1 + K \left(\mathfrak{S}_A - \frac{\mathfrak{S}_B}{K} \right)^2 e^{-\lambda\tau} \right]$$

where $K = k_+ / k_-$, $\lambda = k_+ + k_-$, and \mathfrak{S} is the fractional intensity of state A or B

$$G_c(\tau) = G_D(\tau, N_C, \tau_D)$$

Diffusion term drops out of the ratio

$$\frac{G_b(\tau)}{G_c(\tau)} = \frac{G_b(0)}{G_c(0)} \left(1 + \frac{1}{K} \exp(-\lambda\tau) \right)$$

$$\lambda = k_+ + k_-$$

$$1/\lambda = 24.2 \mu\text{s}$$



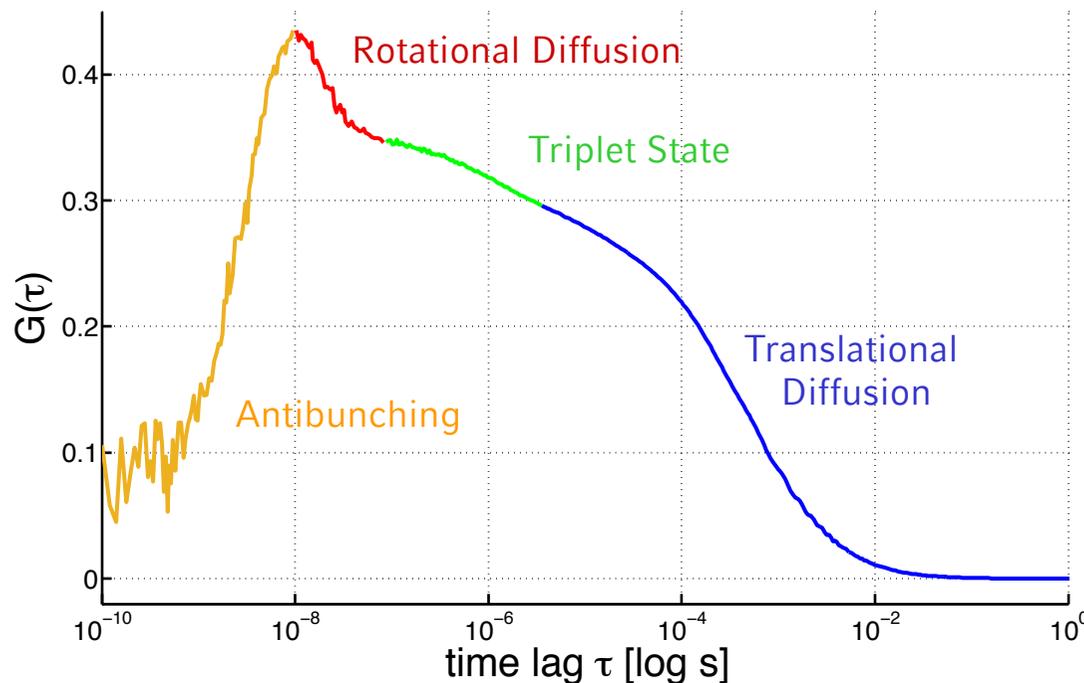
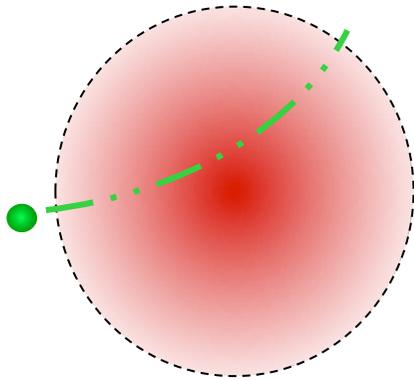
Fluctuation Correlation Spectroscopy (FCS)



Freely diffusing, non-interacting particles in an open volume.

Photons are not detected stochastically, but in bursts when a molecule transverses the probe volume.

Several processes other than diffusion can lead to fluctuations in fluorescence intensity



We can determine:

- Excited state lifetime
- Rotational Diffusion Constant
- Reaction Kinetics
- Triplet-State Lifetime
- Triplet-State Amplitude
- Translational Diffusion Constant
- Concentration



ACF with Multiple Species



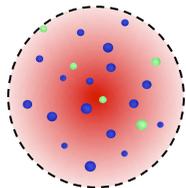
$$G(\tau) = \mathfrak{S}_1^2 G_{D1}(\tau, N_1, \tau_{D1}) + \mathfrak{S}_2^2 G_{D2}(\tau, N_2, \tau_{D2}) \quad \text{2 species}$$

$$\mathfrak{S}_i = \varepsilon_i \langle N_i \rangle / (\varepsilon_1 \langle N_1 \rangle + \varepsilon_2 \langle N_2 \rangle) \quad \text{Fractional Intensity}$$

$$G(\tau) = \sum_{i=1}^M \mathfrak{S}_i^2 G_{Di}(\tau, N_i, \tau_{Di}) \quad \text{Multiple species}$$

FCS measurements in a fluorescent background

With Background

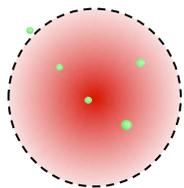


Situation: Large number of weakly fluorescing particles:

i.e. Although $Q_S \gg Q_B$, $N_S \ll N_B \Rightarrow$

$$G_B(0) = \left(\frac{\gamma}{\langle N_B \rangle} \right) \ll \left(\frac{\gamma}{\langle N_S \rangle} \right) = G_S(0)$$

Without Background



$$G(\tau)_{eff} = \mathfrak{S}_S^2 G_{Diff}(\tau, N_S, \tau_{D_S}) + \mathfrak{S}_B^2 G_{Diff}(\tau, N_B, \tau_{D_B})$$

$$G(\tau)_{eff} = \mathfrak{S}_S^2 G_{Diff}(\tau, N_S, \tau_{D_S})$$

0

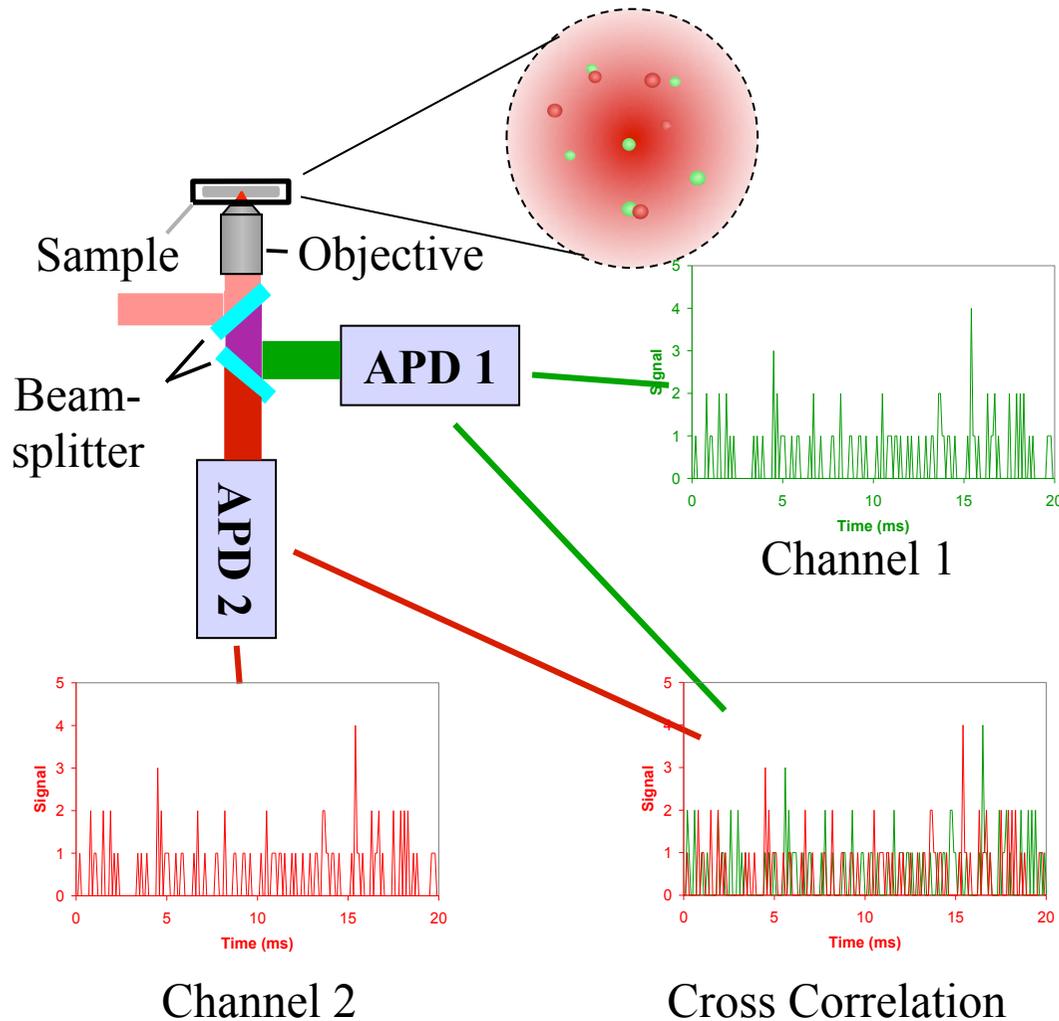
The amplitude of the ACF is reduced by the *square* of the fractional intensity



Fluorescence Cross-Correlation Spectroscopy



Two-Channel Measurements



Type of Experiment	Optics and Channels
Fast Correlation	50/50 Beamsplitter
	Ch1: 50% of original Signal
	Ch2: 50% of original Signal
Two Color Cross-Correlation	Dichroic Mirror
	Ch1: Green
	Ch2: Red
Rotational Diffusion	Polarizing Beamsplitter
	Ch1: \perp Polarized light
	Ch2: \parallel Polarized light

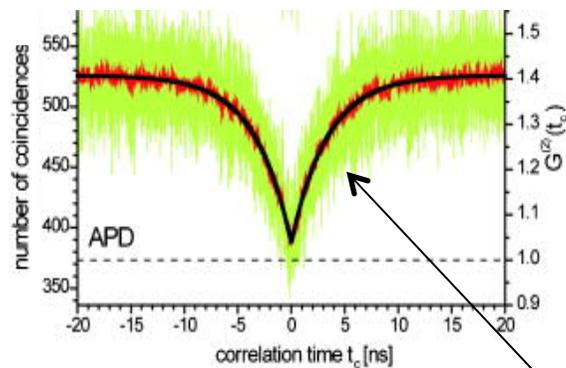


Fast correlation – correlation curves

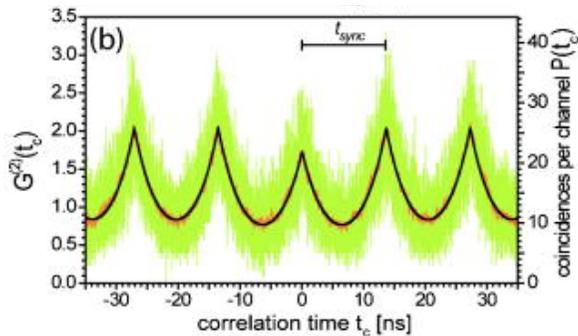


Short-time correlation curves of Rhodamine 110 aqueous solutions measured with APD detectors:

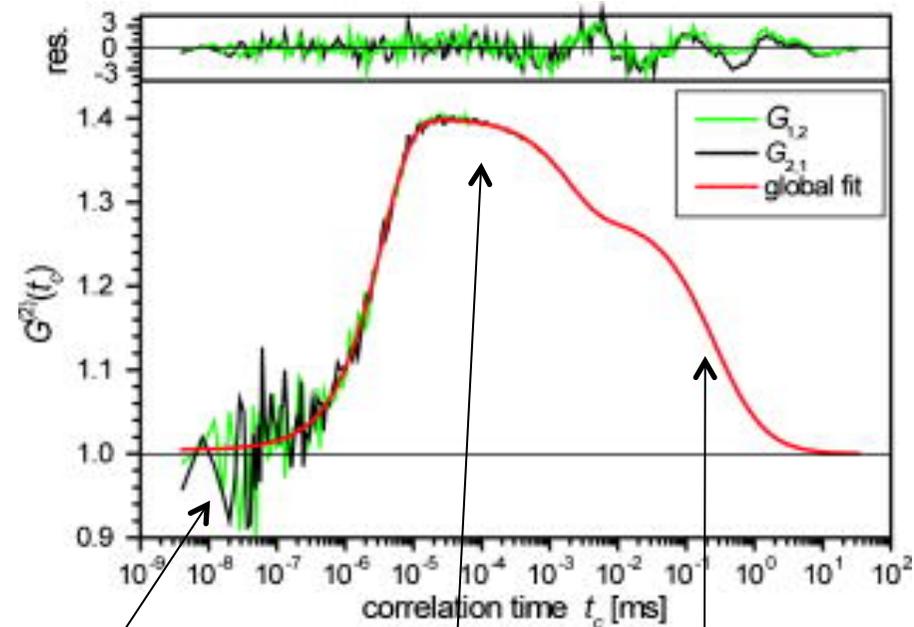
...using cw excitation:



...using pulsed excitation



Full ACF



Antibunching
(No Afterpulsing)

Triplet formation

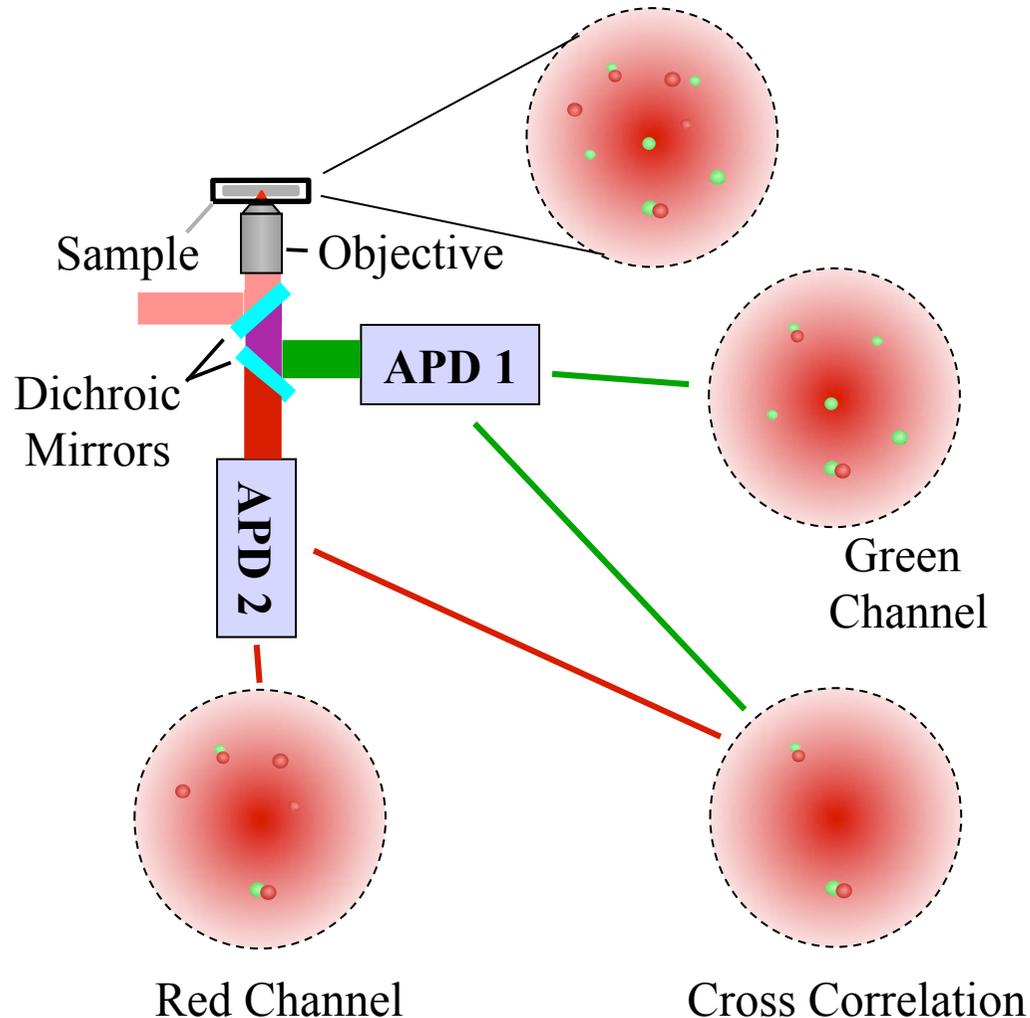
Lateral diffusion



Fluorescence Cross-Correlation Spectroscopy



Two-Channel Measurements



The sample consists of three species:

N_G Particles/complexes with a green label only

N_R Particles/complexes with a red label only

N_{GR} Particles/complexes containing both green and red labels

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

Ideally, only the N_{GR} particles cross correlate

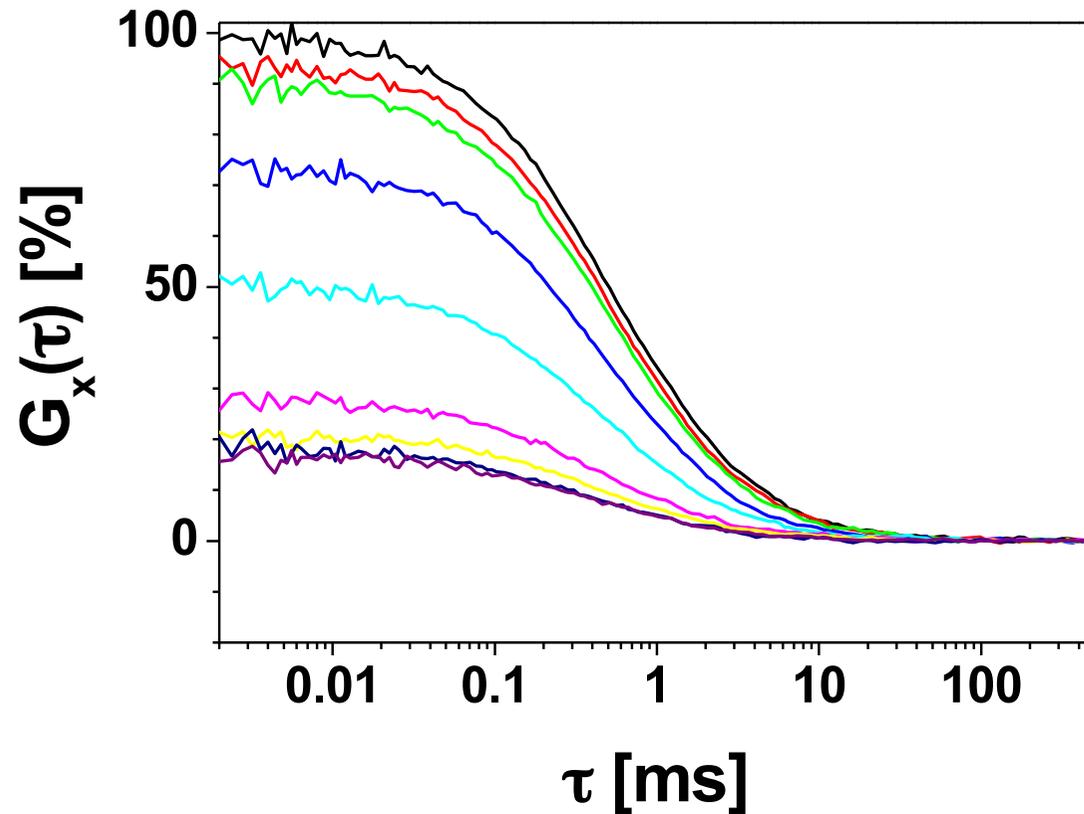
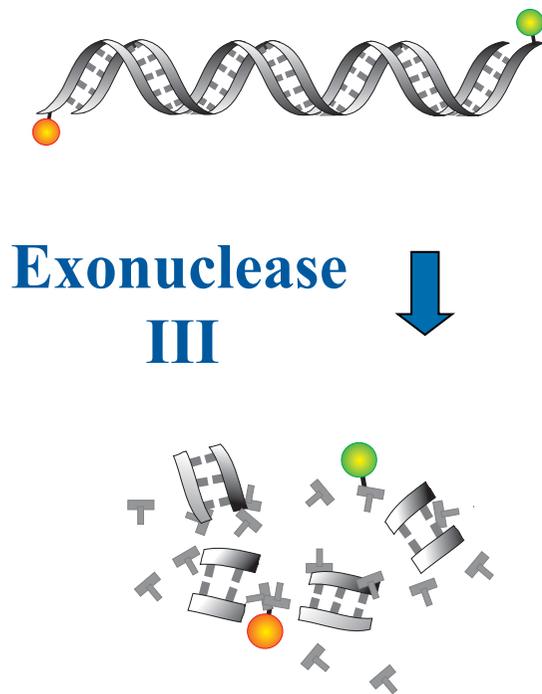
$$G_{GR}(\tau) = \frac{\gamma \langle N_{GR} \rangle G_D(1, D_{GR}, \tau)}{\langle N_G + N_{GR} \rangle \langle N_R + N_{GR} \rangle}$$



Kinetics of DNA Degradation



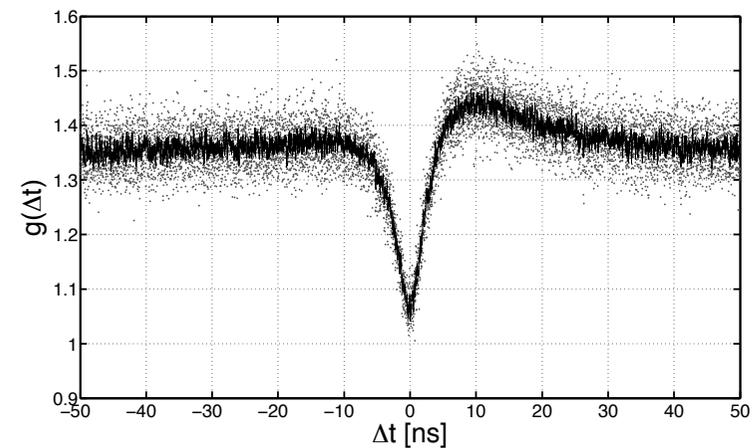
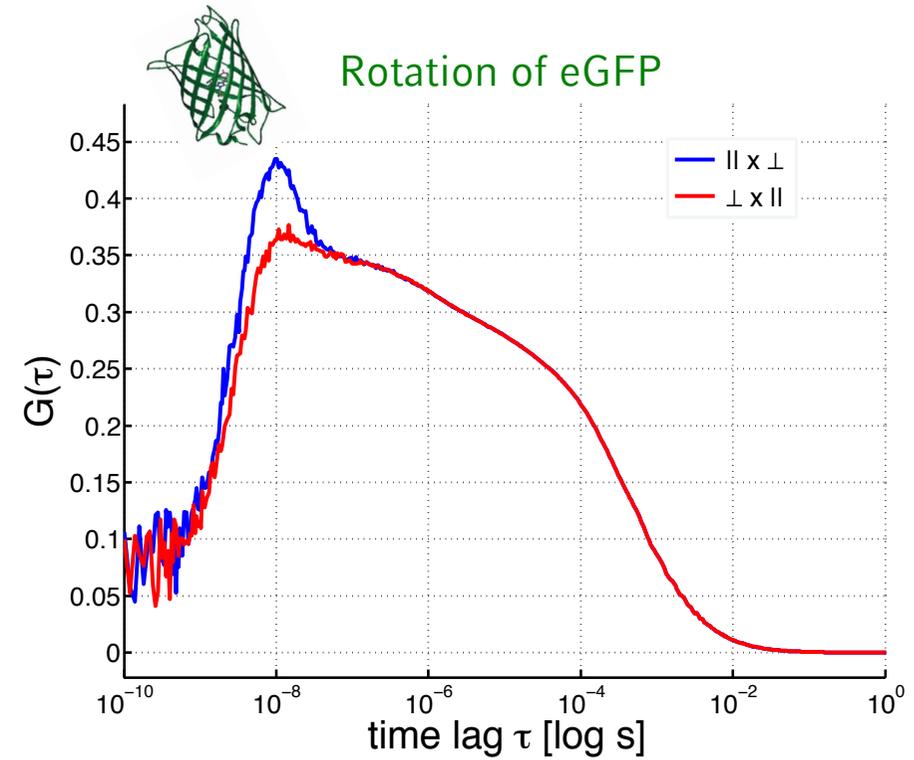
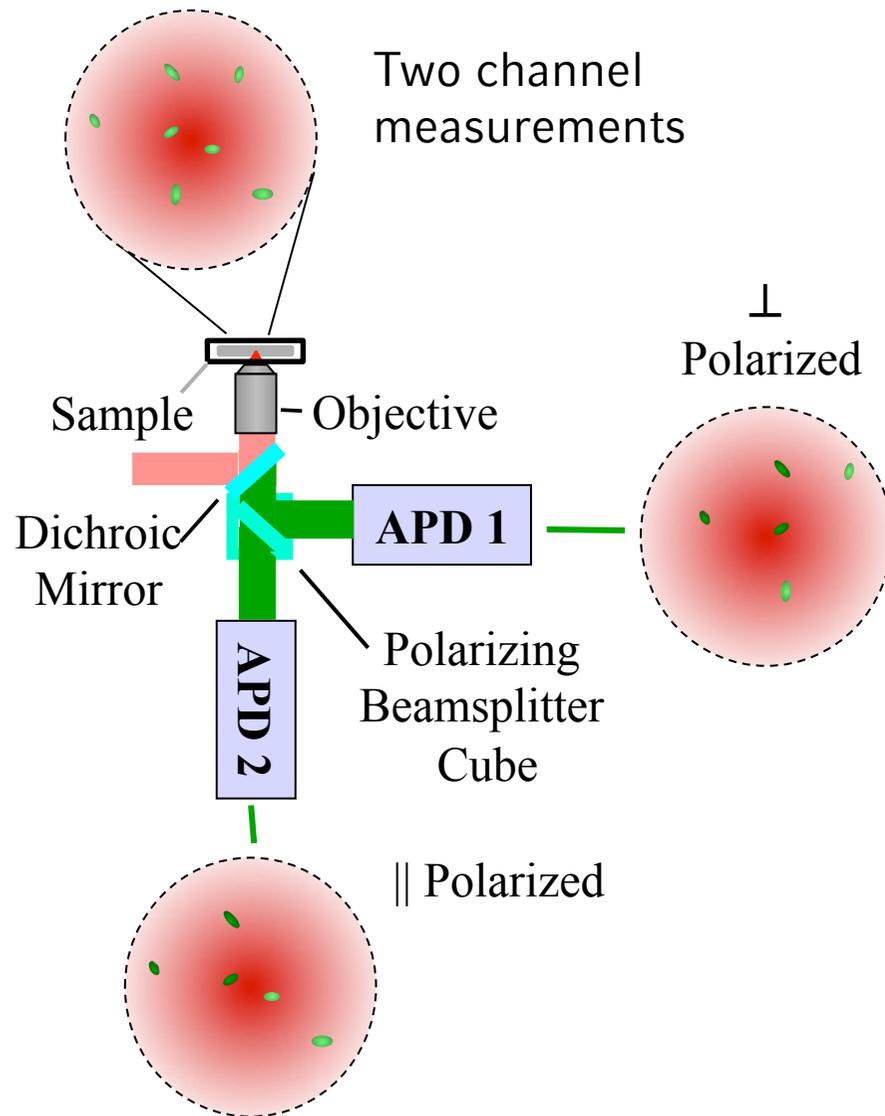
$$G_{GR}(0) = \frac{\gamma \langle N_{GR} \rangle}{\langle N_G + N_{GR} \rangle \langle N_R + N_{GR} \rangle}$$



Ketting, Koltermann, Schwille, Eigen *PNAS* (1998) 95:1416



Rotational Diffusion





Fluctuations in Space

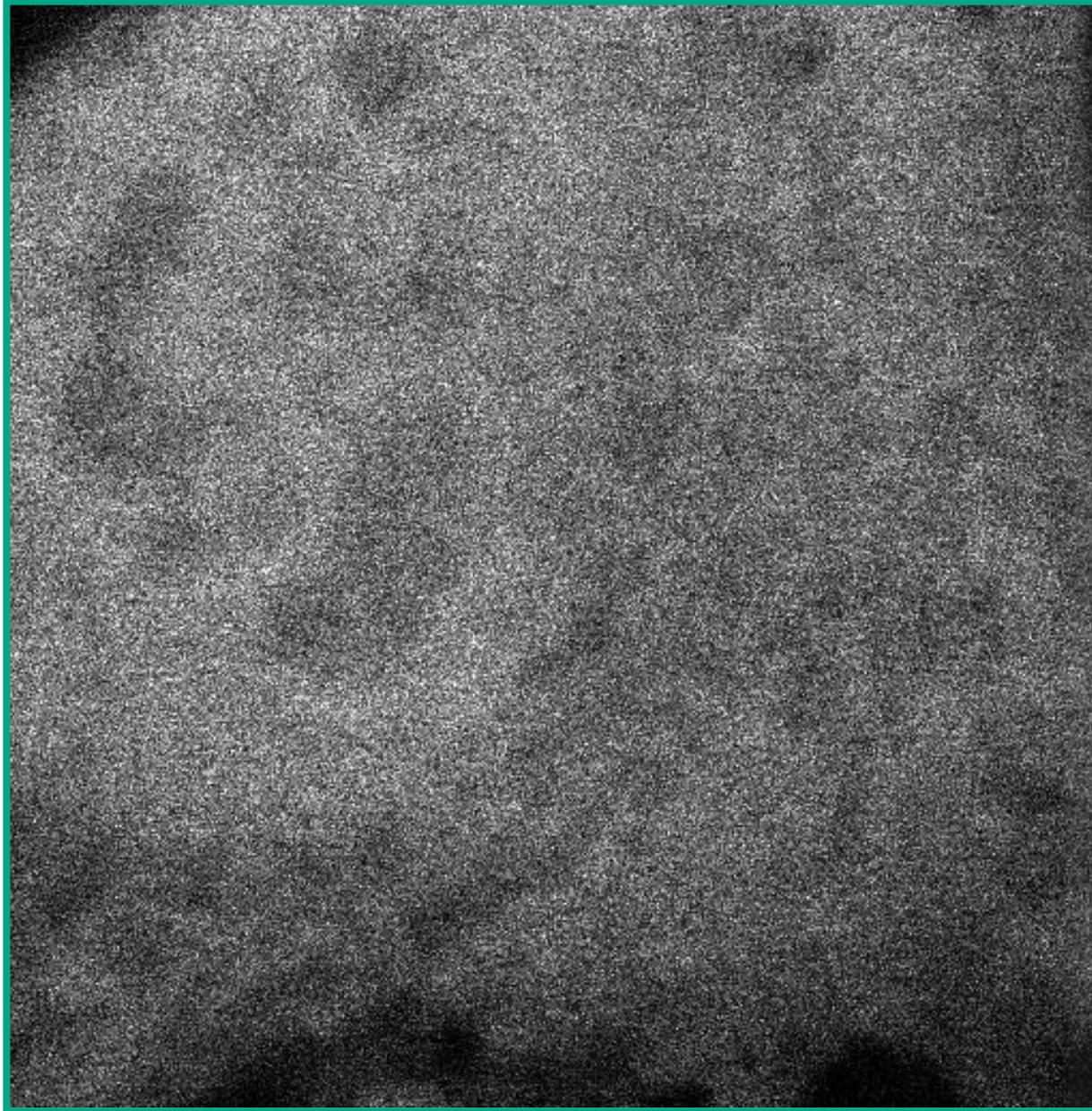
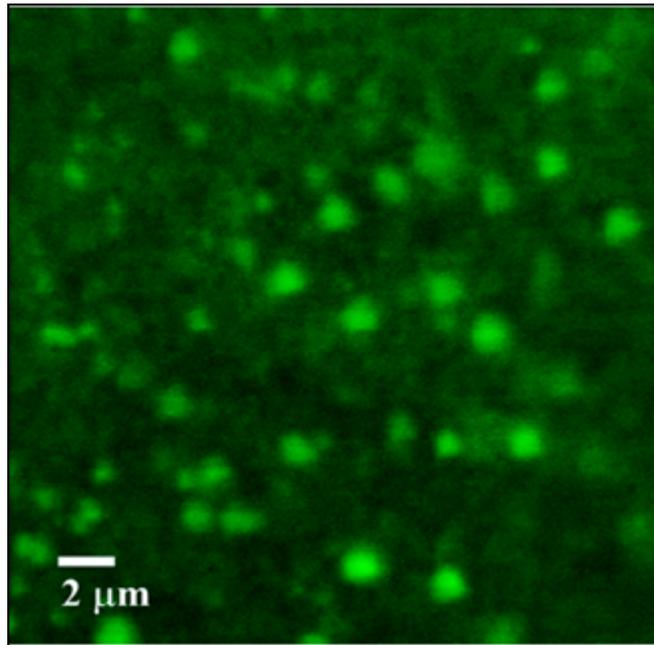


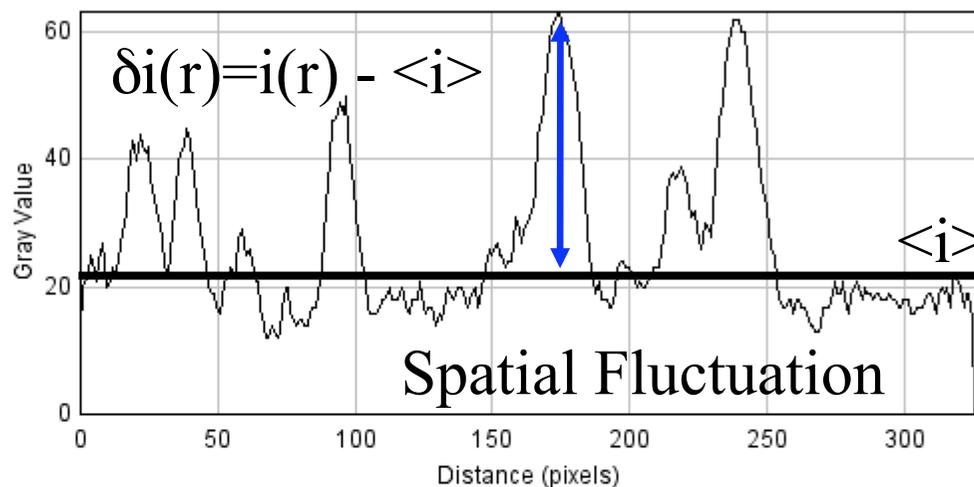
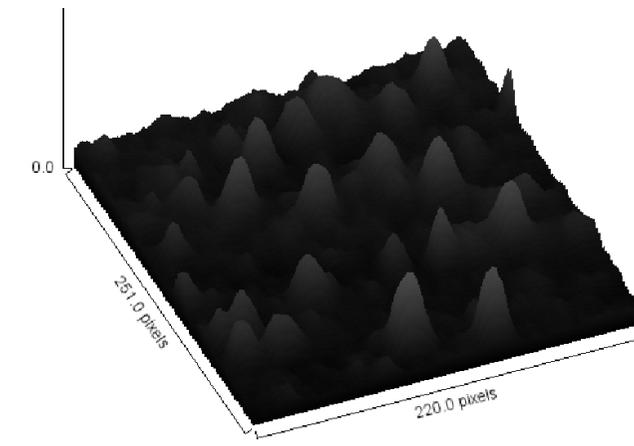


Image as a Record of Spatial Fluctuations



Ergodic: i.e. every sizeable sampling of the process is representative of the whole

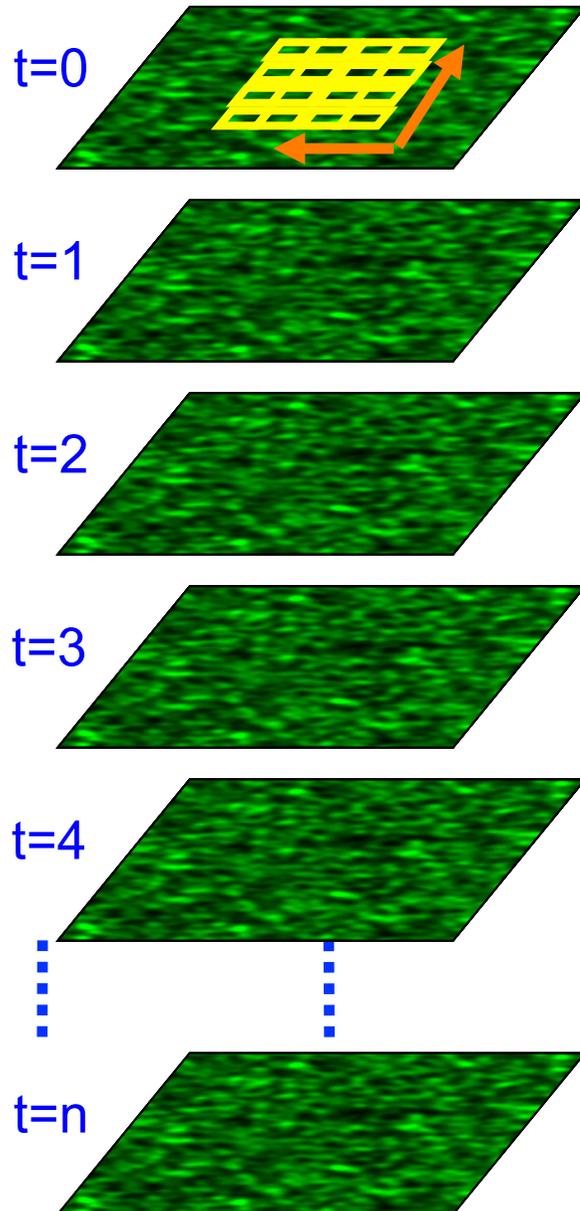
the time average is equal to the ensemble average



The Image records fluorescence fluctuations in space which reflect the distribution of labeled molecules

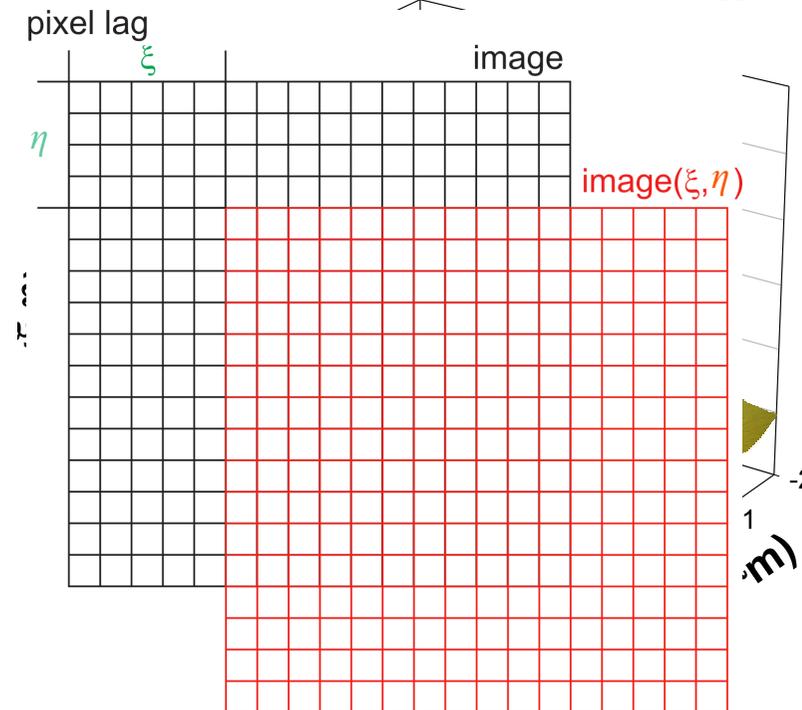


Image Correlation Spectroscopy (ICS)



Spatial autocorrelation
of $\delta i(x,y,t) = i(x,y,t) - \langle i \rangle$
for each image in x, y

$$r_{11}(0,0)_j = \frac{1}{\langle N \rangle}$$



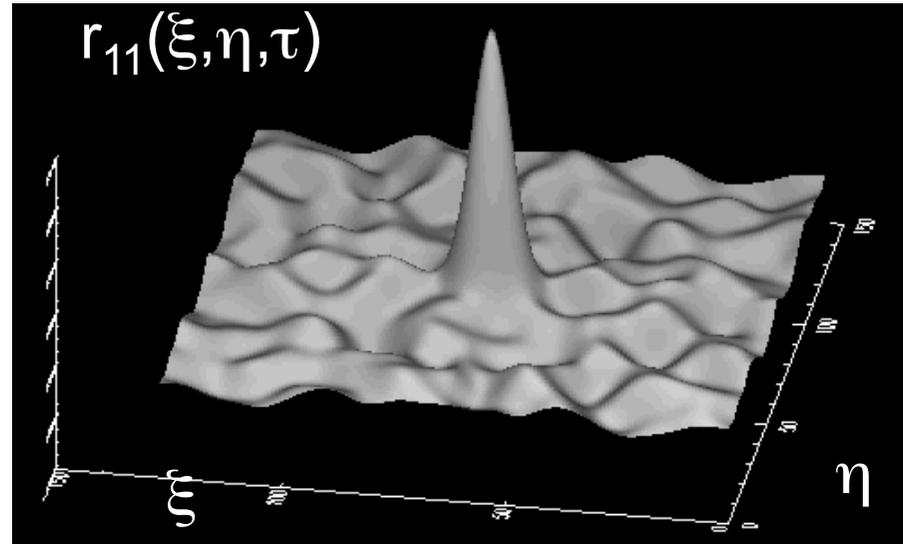
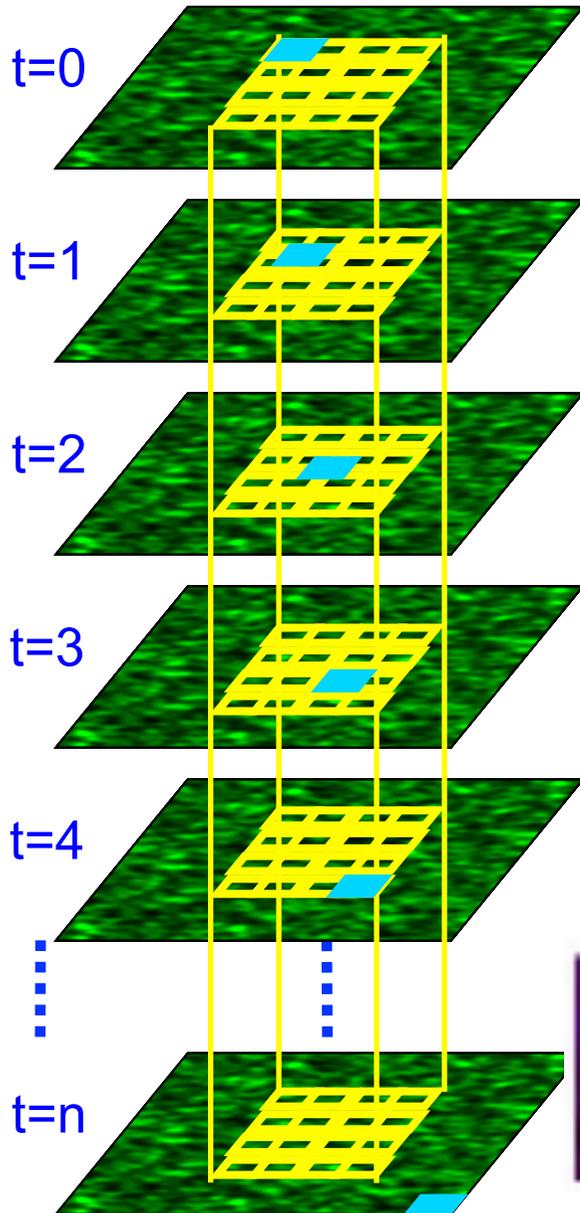
$$r_{11}(\xi, \eta)_j = \frac{\langle \delta i(x + \xi, y + \eta) \delta i(x, y) \rangle}{\langle i \rangle^2}$$

Petersen *et al.* *Biophys. J.* 65, 1135-1146 (1993);
Wiseman and Petersen, *Biophys. J.* 76, 963-977 (1999)



Spatio-Temporal Image Correlation Spectroscopy

Spatial correlation
as a function of time



$$r_{11}(\xi, \eta, \tau) = \frac{\langle \delta i(x, y, t) \delta i(x + \xi, y + \eta, t + \tau) \rangle}{\langle i \rangle_t \langle i \rangle_{t+\tau}}$$

Hebert et al. Biophys. J. 88-3601 (2005)

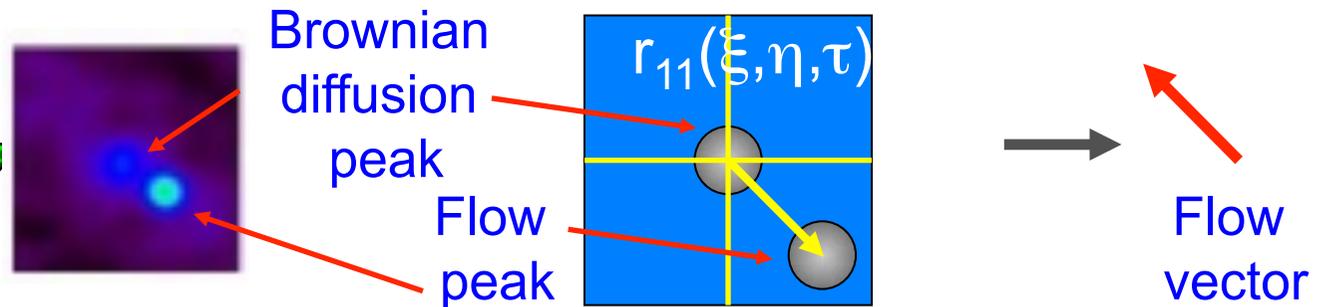


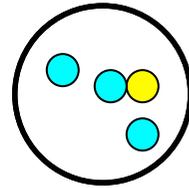
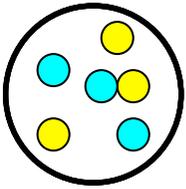


Image Cross-Correlation Spectroscopy (ICCS)

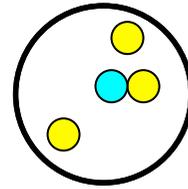


Wiseman et al., *J. Microscopy* 200, 14-25 (2000)

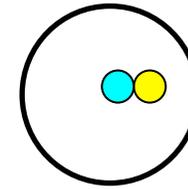
Biological
Sample



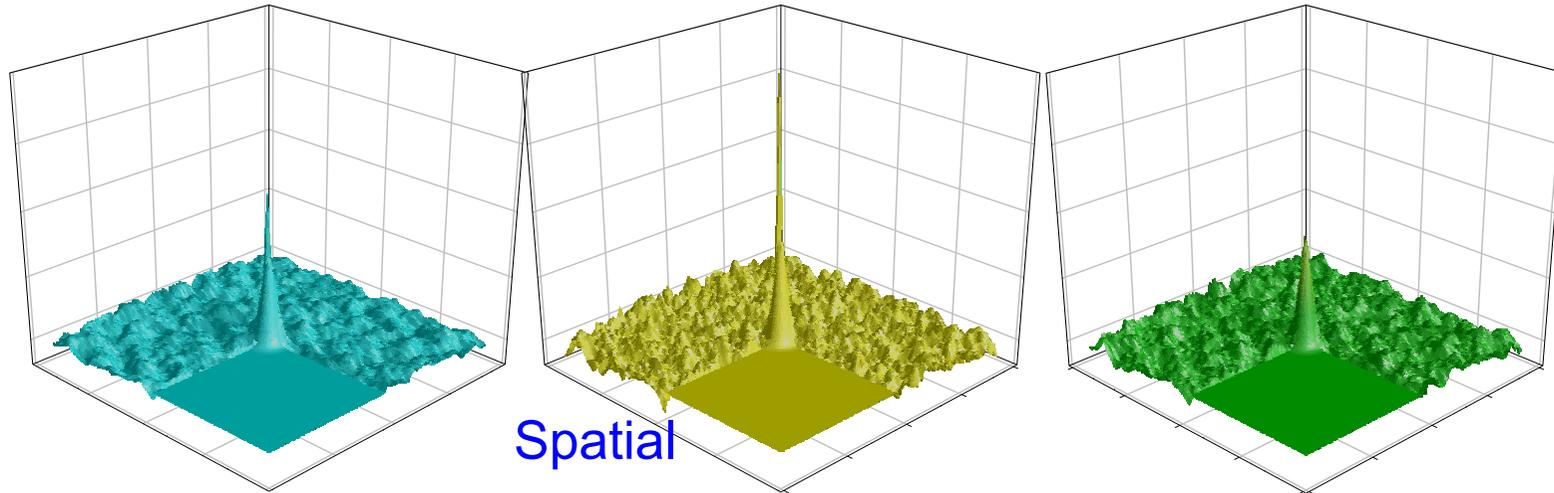
Auto1



Auto2



Cross



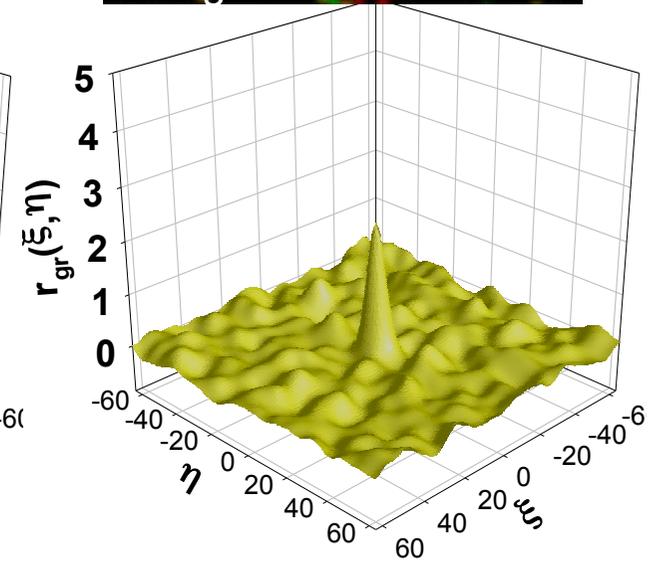
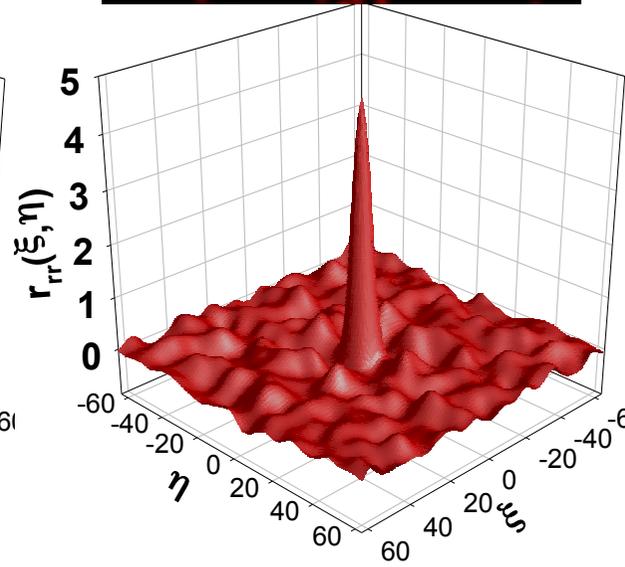
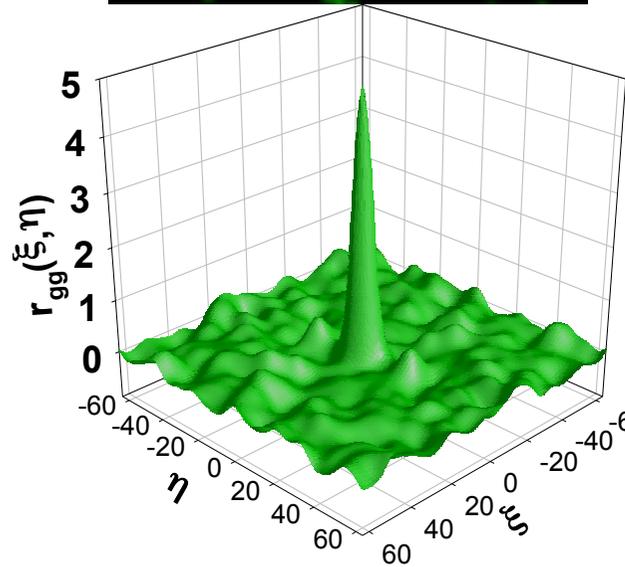
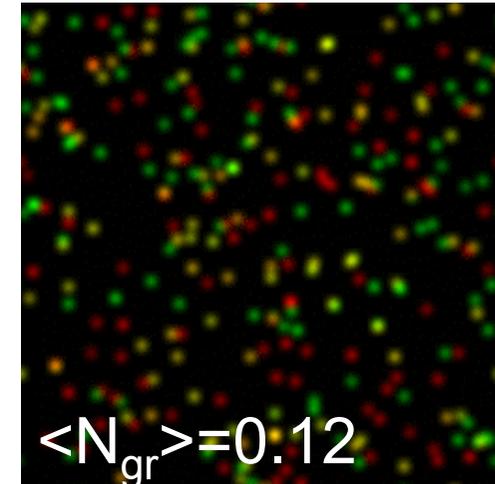
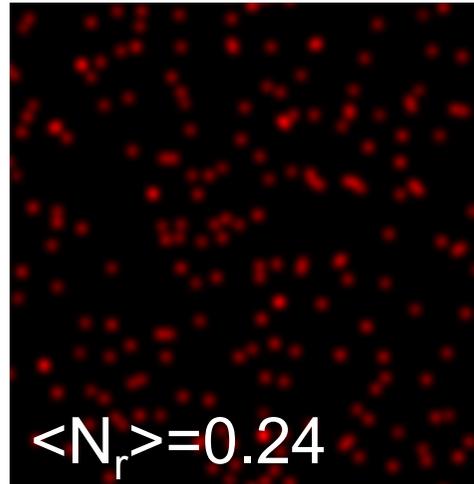
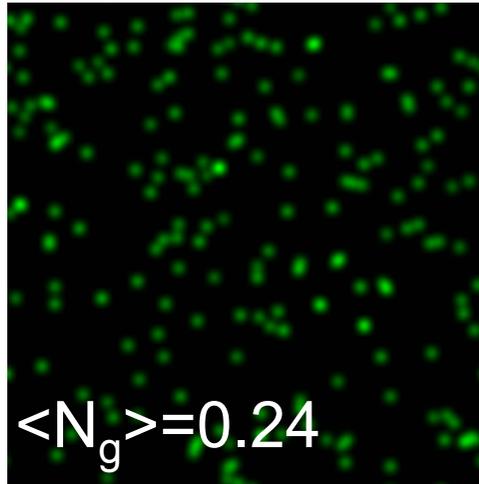
Spatial

Correlation

$$r_{ab}(\xi, \eta, 0)_j = \frac{\langle \delta i_a(x, y, t) \delta i_b(x + \xi, y + \eta, t) \rangle}{\langle i_a \rangle_t \langle i_b \rangle_t}$$



Spatial ICCS Simulations



$$\langle N_{gr} \rangle = \frac{g_{gr}(0,0)}{g_g(0,0) g_r(0,0)} = 0.10$$



ICS (Image Correlation Spectroscopy) spatial autocorrelation of an image

Size and number of membrane aggregates (Paul Wiseman & Nils Petersen)

tICS (Temporal Image Correlation Spectroscopy)

time autocorrelation at one pixel (Mamta Srivastava Paul Wiseman, Nils Petersen)

STICS (Spatio-Temporal Image Correlation Spectroscopy)

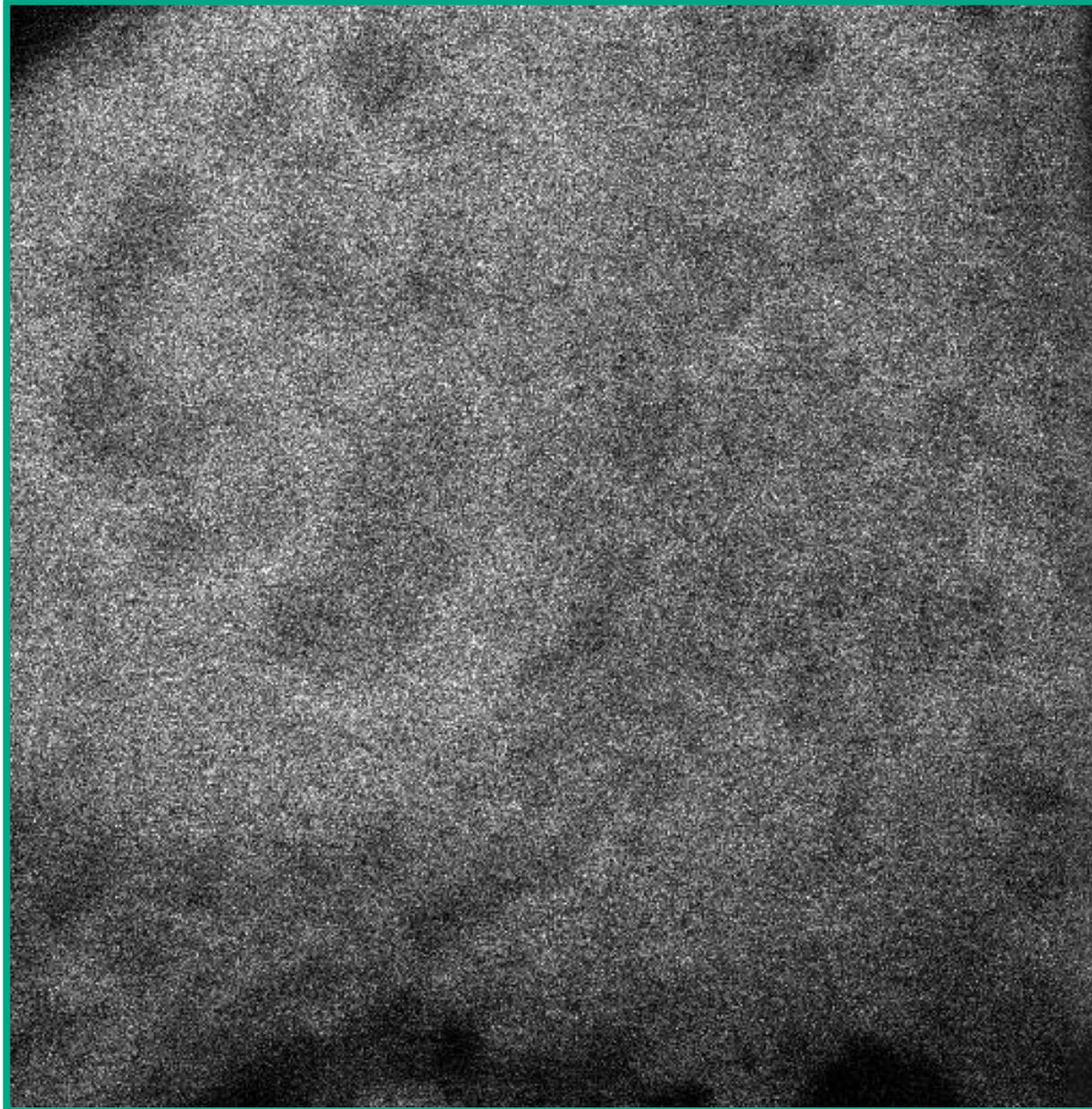
Diffusion and velocity in 2-D (Ben Herbert & Paul Wiseman)

kICS (k-space Image Correlation Spectroscopy)

Distinguish between diffusion and binding on the basis of spatial correlations
(Paul Wiseman, David Kolin, David Ronis)

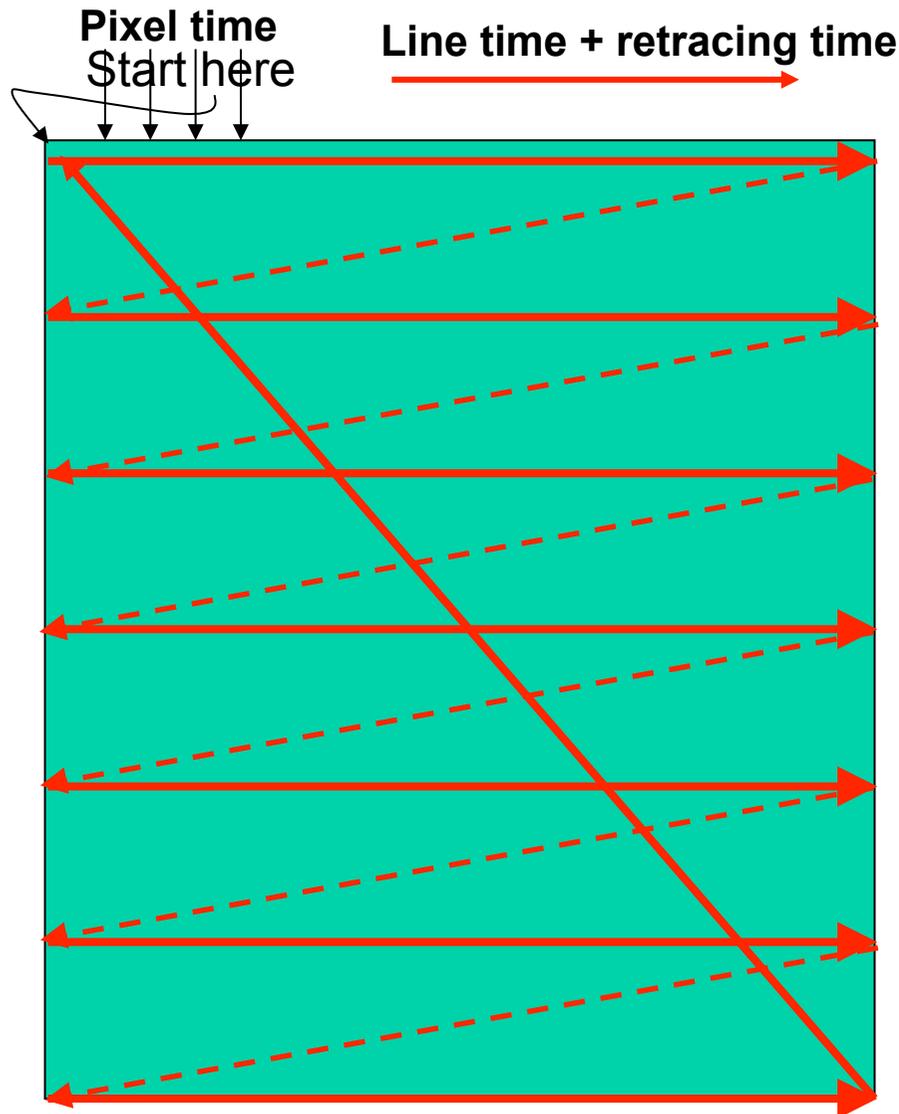


Fluctuations in Time and Space: RICS

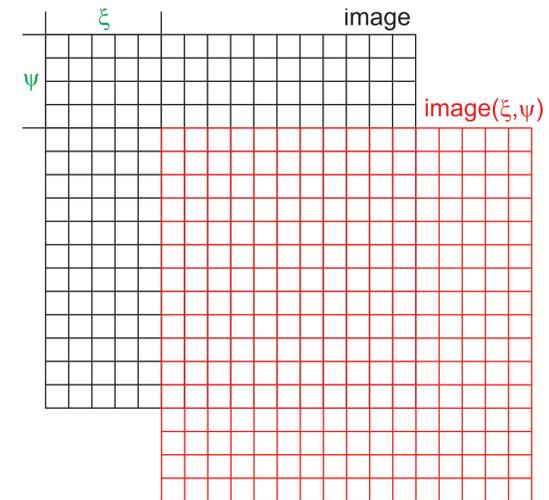




Raster Image Correlation Spectroscopy



$$G_{ICS}(\xi, \psi) = \frac{\langle I(x, y)I(x + \xi, y + \psi) \rangle_{XY}}{\langle I(x, y) \rangle_{XY}^2}$$



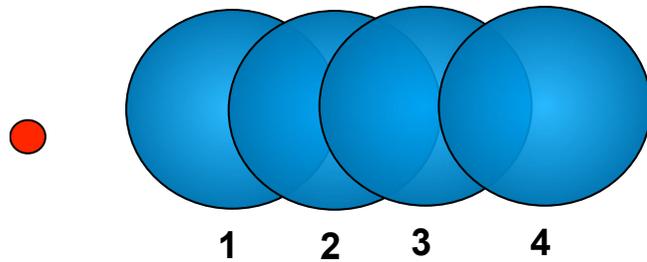
$$G_{RICS}(\xi, \psi) = \frac{\gamma}{N} \left(1 + \frac{4D(\tau_p \xi + \tau_l \psi)}{w_0^2} \right)^{-1}$$

$$\left(1 + \frac{4D(\tau_p \xi + \tau_l \psi)}{w_z^2} \right)^{-1/2} \exp \left(-\frac{\delta r^2 (\xi^2 + \psi^2)}{w_0^2 + 4D(\tau_p \xi + \tau_l \psi)} \right)$$

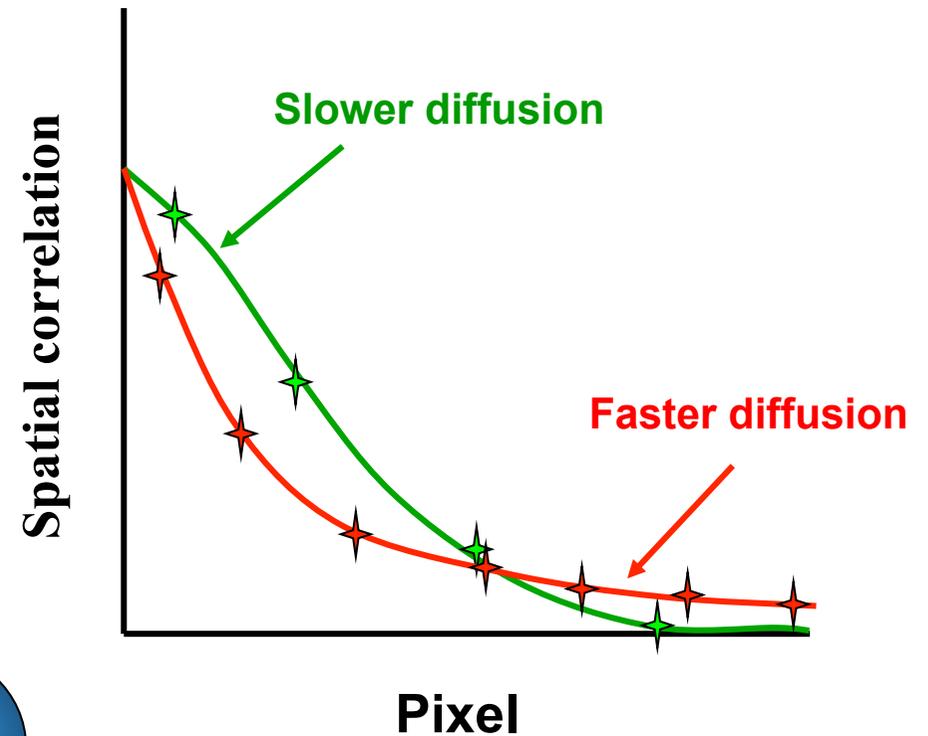
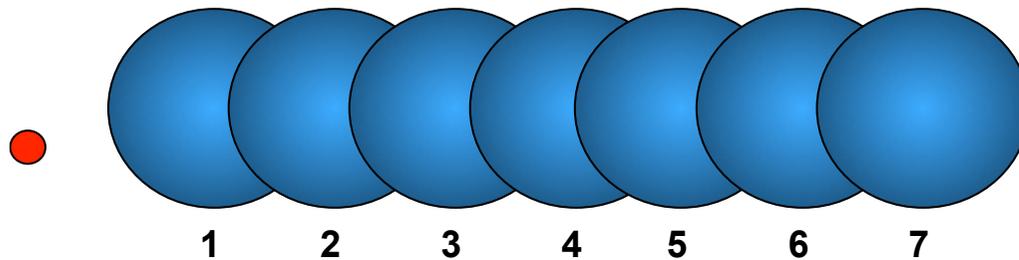


Temporal information hidden in the raster-scan image:

Situation 1: slow diffusion



Situation 2: fast diffusion

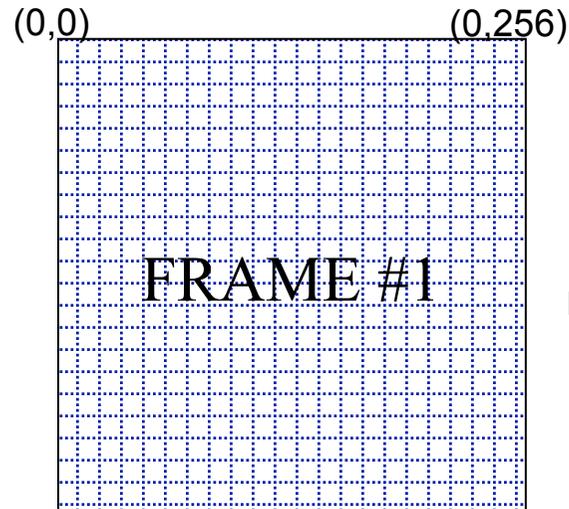




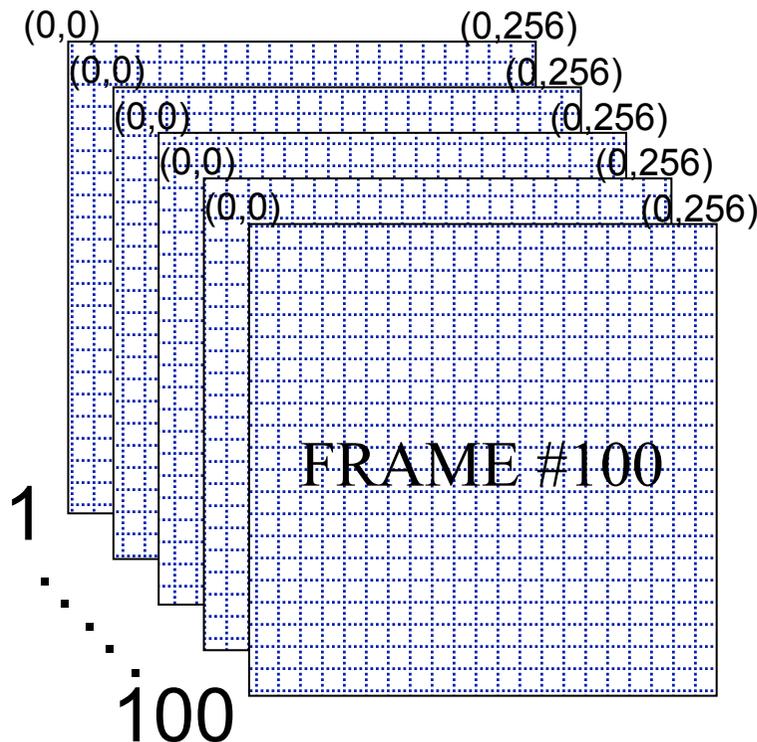
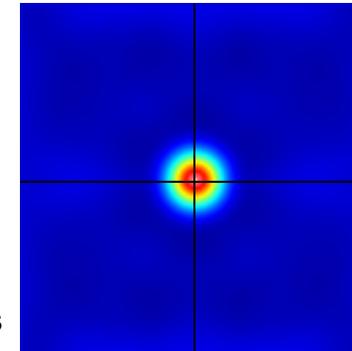
Averaging in RICS



The image correlation is performed within each frame



Spatially correlate each frame
Individually then take the average of all the frames

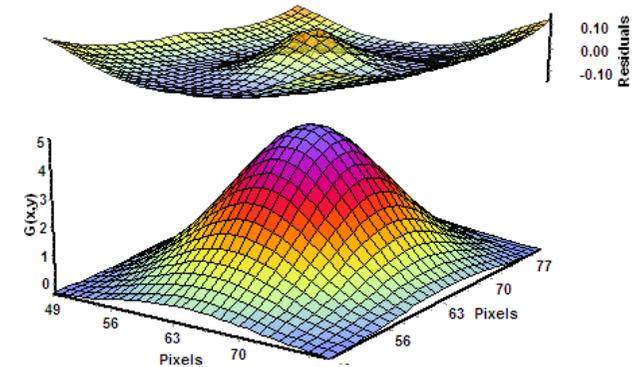
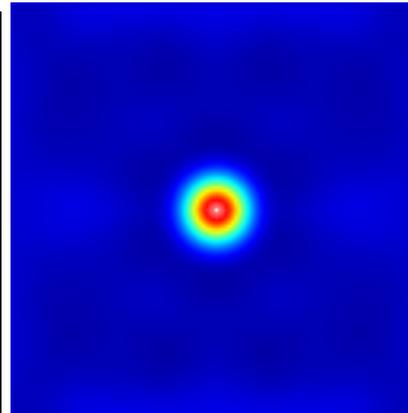
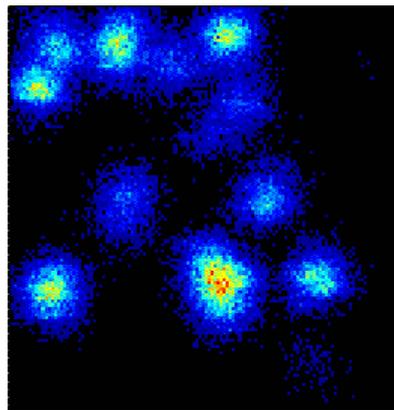




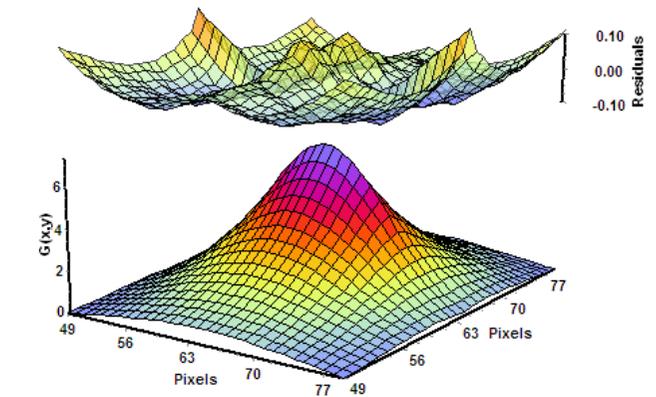
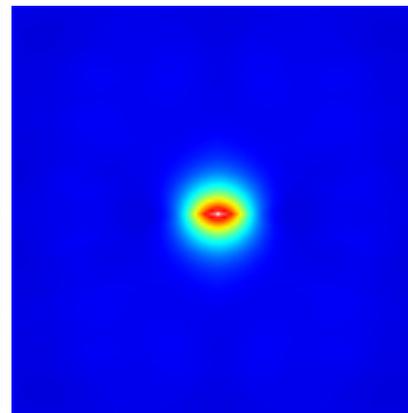
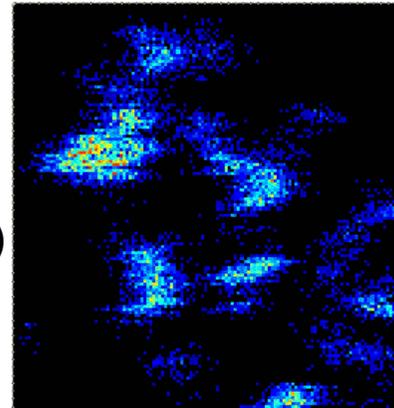
RICS Simulations for Different Diffusion Coefficients



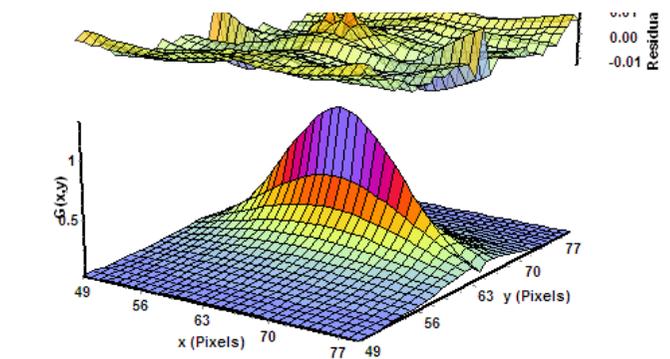
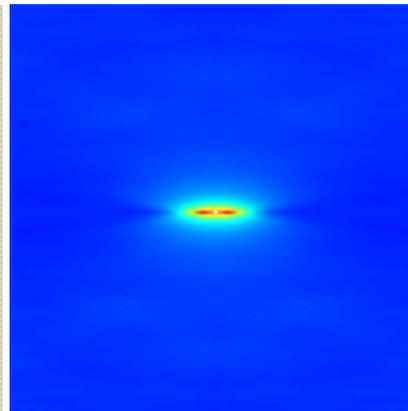
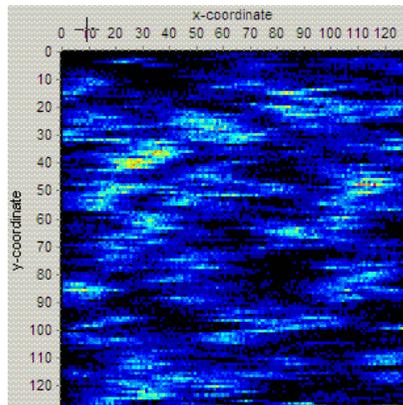
$D = 0.1 \mu\text{m}^2/\text{s}$
(membrane proteins)



$D = 5.0 \mu\text{m}^2/\text{s}$
(40 nm beads)



$D = 90 \mu\text{m}^2/\text{s}$
(EGFP)

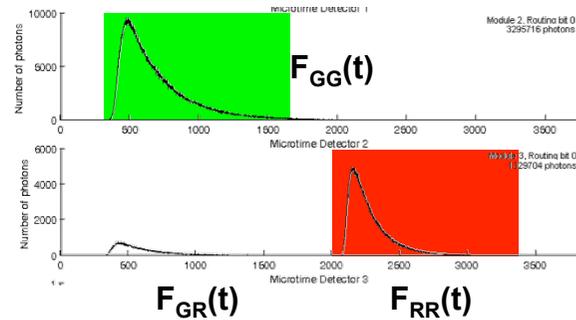




Cross-Correlation-Free RICS



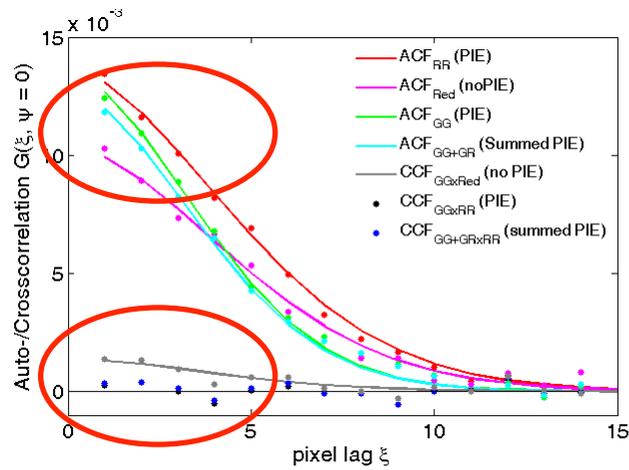
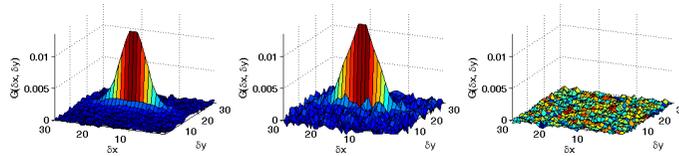
eGFP + mCherry in HeLa cells



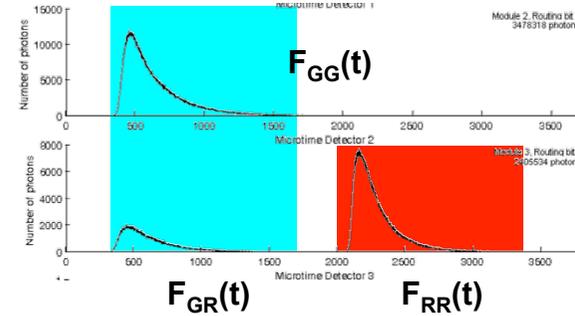
Green APD

Red APD

ACF eGFP ACF mCherry CCF

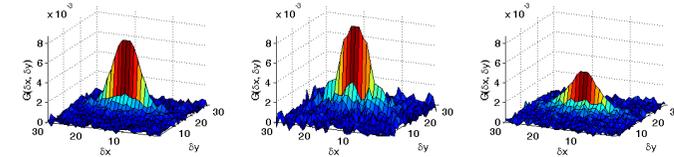


eGFP-mCherry in HeLa cells



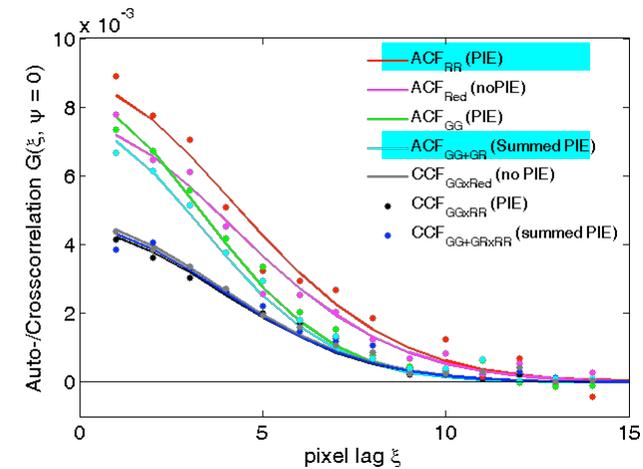
3D RICS

ACF eGFP ACF mCherry CCF



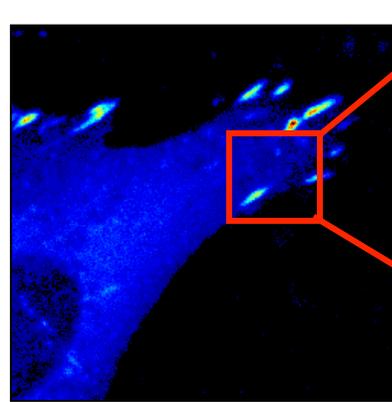
3D RICS

$G(\xi, \psi=0)$

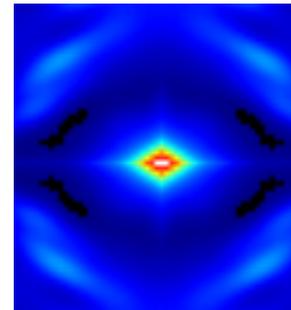
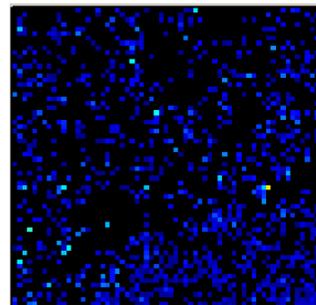
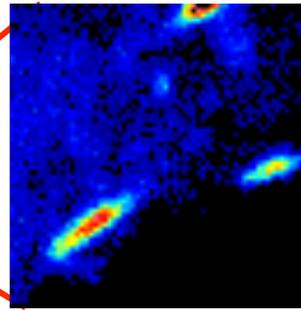




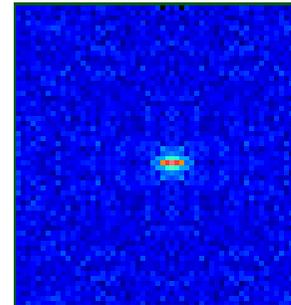
Removal of Immobile Structures and Slow Moving Features



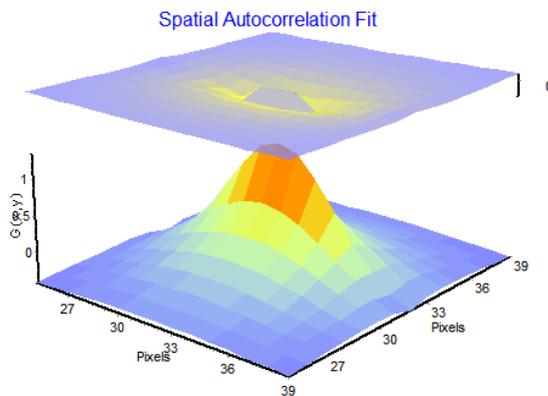
What is left after removal



Spatial ACF
No removal



Spatial ACF
With removal



Fit using 3-D diffusion formula

Pixel size = $0.092 \mu\text{m}$

Pixel time = $8 \mu\text{s}$

Line time = 3.152ms

$W_0 = 0.35 \mu\text{m}$

$G_1(0) = 0.0062$

$D_1 = 7.4 \mu\text{m}^2/\text{s}$

$G_2(0) = 0.00023$

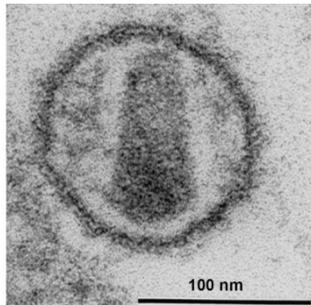
$D_2 = 0.54 \mu\text{m}^2/\text{s}$

$\text{Bkgd} = -0.00115$

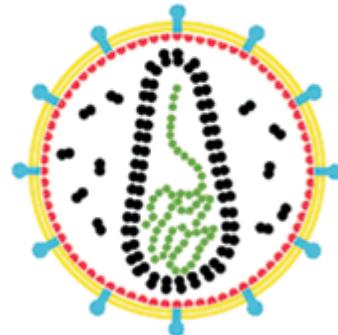
HIV: membrane enveloped retrovirus with RNA genome

Electron Microscopy Images of HIV

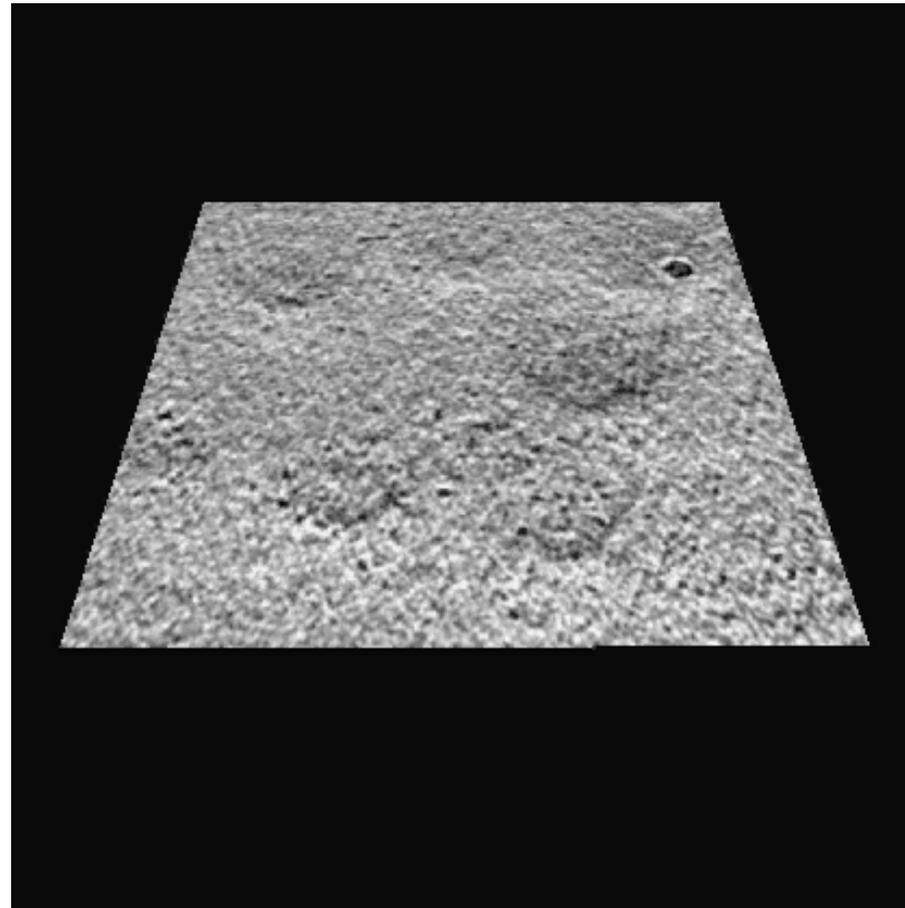
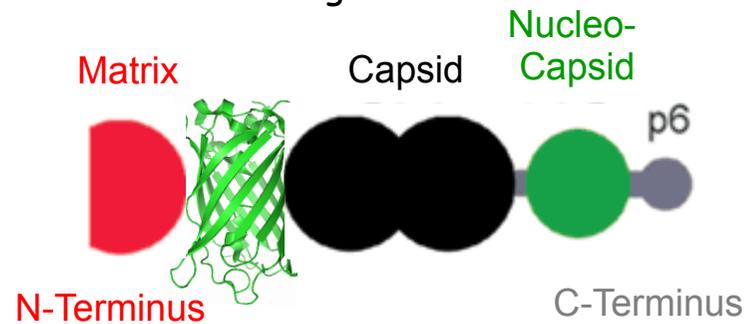
Welker, et al. 2000 J Virol **74**:1168.



Briggs et al. 2004 Nature Struct Mol Biol **11**:672



GFP-Gag Protein



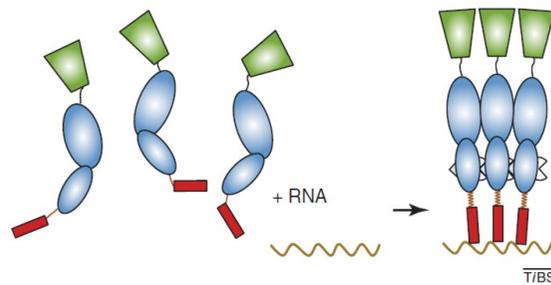
Briggs et al. 2006 Structure **14**:15
Supplemental Movie



Contributions of Membrane, RNA and CA-CA Interactions



- PI(4,5)P₂ induces conformational switch in Gag,
Myristate gets exposed and binds PM Ono et al.(2004), PNAS **101**, 14889-14894
- Matrix domain G2A mutant Göttinger et al. (1989), PNAS **86**, 5781-5785
 - Defective for assembly and particle release
 - Interacts much less with membranes!
- In vitro: Gag assembles into VLPs upon addition of nucleic acid
Campbell et al.(1995), J Virology **69**, 6487-6497
 - RNA primes Gag for interactions ?



from Rein et al. (2011), TIBS, 1-8

- Targeting Gag away from RNA or Gag
 - NC C15A inhibits Gag-RNA interaction
Poon et al. (1996), J Virology **70**, 6607-6616
 - CA W184A/M185A inhibits CA-CA
von Schwedler et al. (2003), J Virology **77**, 5439-5450

