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# Systematic evaluation of fluorescence correlation spectroscopy data analysis on the nanosecond time scale

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### **Supporting Information**

This material contains six figures S1-S6 with data as described in the figure captions and discussed in more detail within the main text.

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# Figure S1:



**Fig. S1:** Noise amplitudes c(p) determined from a standard FCS sample as function of excitation power p. The data was recorded at 20°C for 15 minutes per data curve. The grey line is a data fit of the function  $c(p)=a*p^m$  with  $a=(0.005\pm0.001) \text{ s}^{1/2}$  and  $m=-0.77\pm0.05$ .

# Figure S2:

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**Fig. S2:** Fit parameter distributions for the fitting results on simulated data sets as described in the main text. (A) Exemplary data curve with noise of  $c=6x10^{-6} s^{1/2}$  overlaid with a function fitted by the LM algorithm as implemented in Mathematica. (B) Fit parameter distributions showing correlations between the two amplitudes A and R (closed squares) as generated from fitting 100 simulated FCS curves in each data set. Four data sets are displayed for data simulated with a noise amplitude of  $c=6x10^{-6} s^{1/2}$  or  $c=15x10^{-6} s^{1/2}$ , A=-2, R=1, k<sub>AB</sub>=0.4, and k<sub>rot</sub>=0.1 (or 0.05 or 0.025). No correlation appears between A<sub>AB</sub> and corresponding  $\chi^2$  values (open circles). N<sub>false</sub> indicates the number of unreasonable data fits with A<sub>AB</sub><-3 (from a total of 100 data fits).

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# Figure S3:

**Fig. S3:** Dynamical fingerprints determined for experimental FCS data of a mixture of freely diffusing fluorophore MR121 (~1 nM) and Trp (10 mM) in PBS buffer with 60% sucrose as presented in Fig. 1. Dynamical fingerprints were determined three times from an identical FCS curve with  $10^5$ ,  $10^7$ , or  $10^9$  iterations (from left to right). The insets show the likelihood for the corresponding series of iteration steps.

# **Figure S4:**



**Fig. S4:** Dynamical fingerprints generated for simulated FCS data. Data was simulated using the model function Eq. 1 overlaid with noise following Eq. 2 with a noise amplitude  $c=9x10^{-6} s^{1/2}$ . Here the starting value for the antibunching time constant was varied ( $10^{-6}$  ms for black,  $9x10^{-7}$  ms for cyan,  $8x10^{-7}$  ms for green,  $7x10^{-7}$  ms for red,  $6x10^{-7}$  ms for pink,  $5x10^{-7}$  ms for blue) while the relaxation time constants were kept constant in all simulations at  $\tau_{ab}=10^{-6}$  ms,  $\tau_{rel}=10^{-5}$  ms,  $\tau_{isc}=3x10^{-4}$  ms (indicated by black lines in the graph). The inset shows all fitted and the simulated data curve.

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**Fig. S5:** Dynamical fingerprints generated for simulated FCS data with a rotational correlation time constant approaching the antibunching time scale. Data was simulated using the model function Eq. 1 overlaid with noise following Eq. 2 with a noise amplitude  $c=9x10^{-6} s^{1/2}$ . The relaxation time of the rotational diffusion correlation decay was varied from  $10^{-5}$  (top) to  $5x10^{-6}$  (bottom) while the relaxation time constants for the antibunching signature and for the decay on larger times was kept constant at  $10^{-6}$  ms and  $3x10^{-4}$  ms, respectively (all indicated by vertical lines in the graphs). The three fingerprints were determined with a slight variation of the antibunching relaxation time starting value ( $1x10^{-6}$  for black;  $8x10^{-7}$  for red,  $6x10^{-7}$  for blue).

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Figure S6:



**Fig. S6:** Dynamical fingerprints determined for experimental FCS data that was recorded for mixtures of freely diffusing fluorophore ATTO655 (~1 nM) and Trp (10 mM) in PBS buffer with various concentrations of sucrose (as indicated in the figure) at 20°C. The fingerprints reveal the same components as shown in Fig. 4 for MR121/Trp with a small additional decay component that mixes with the previous decays and effectively broadens the estimated fingerprint peaks.