

The Effects of High Intensity X-ray Laser Pulses in MHz-XPCS Measurements of Protein Solution Dynamics



Stockholm
University

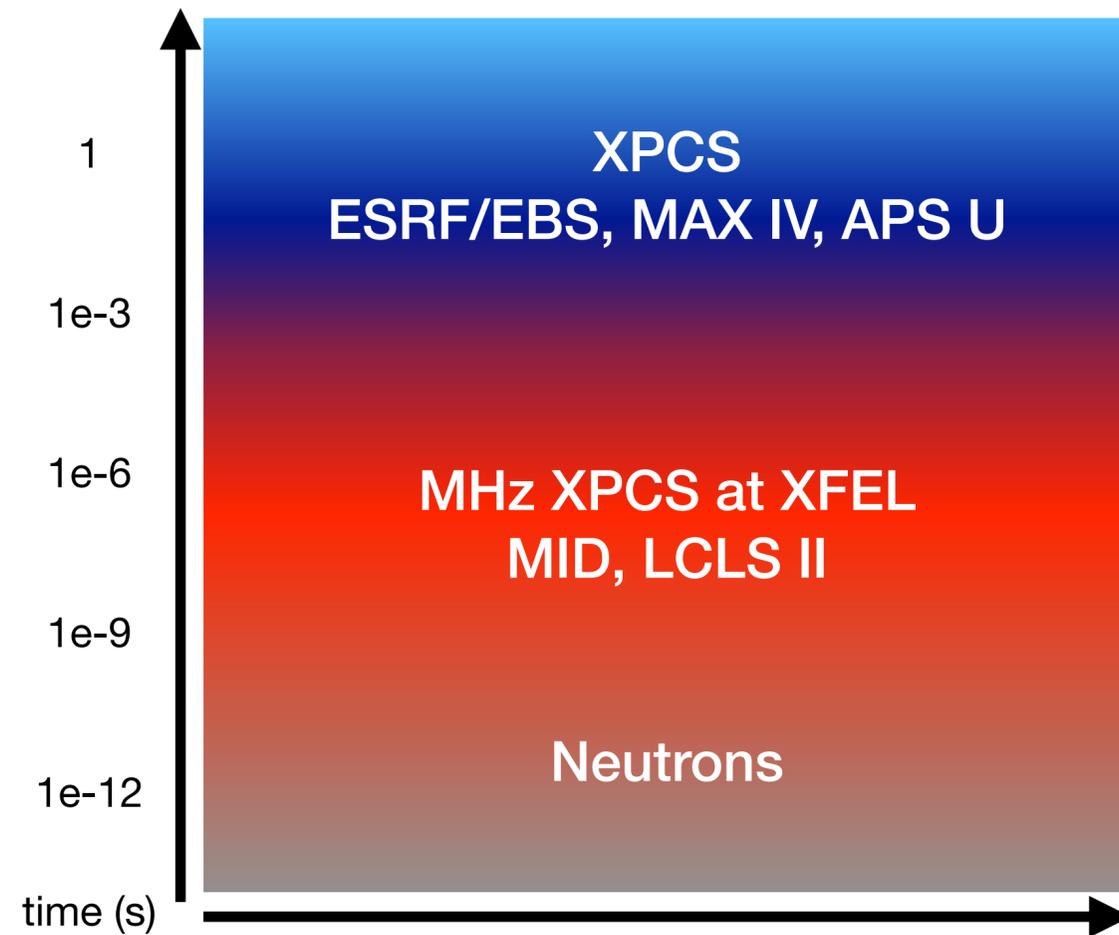


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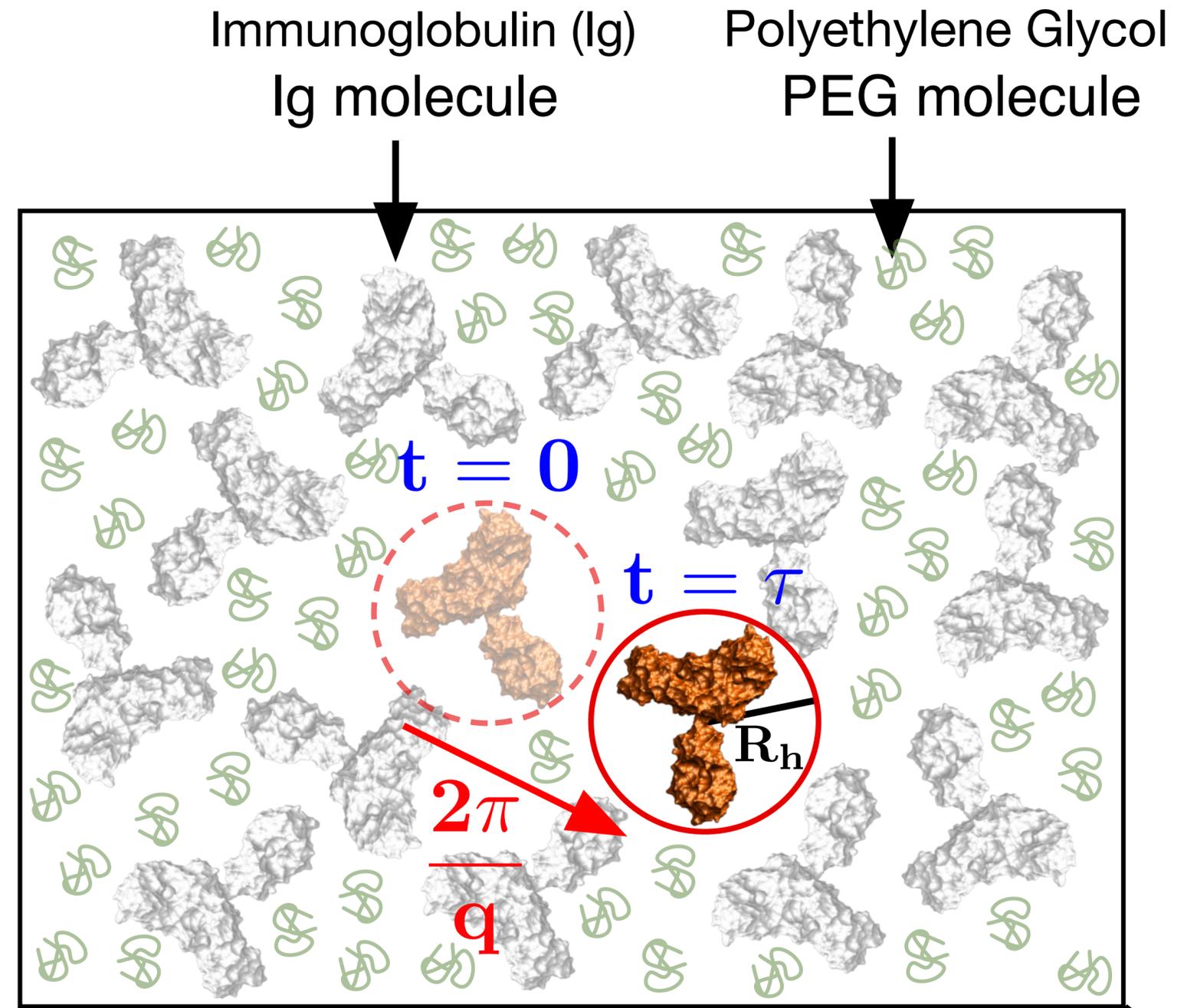
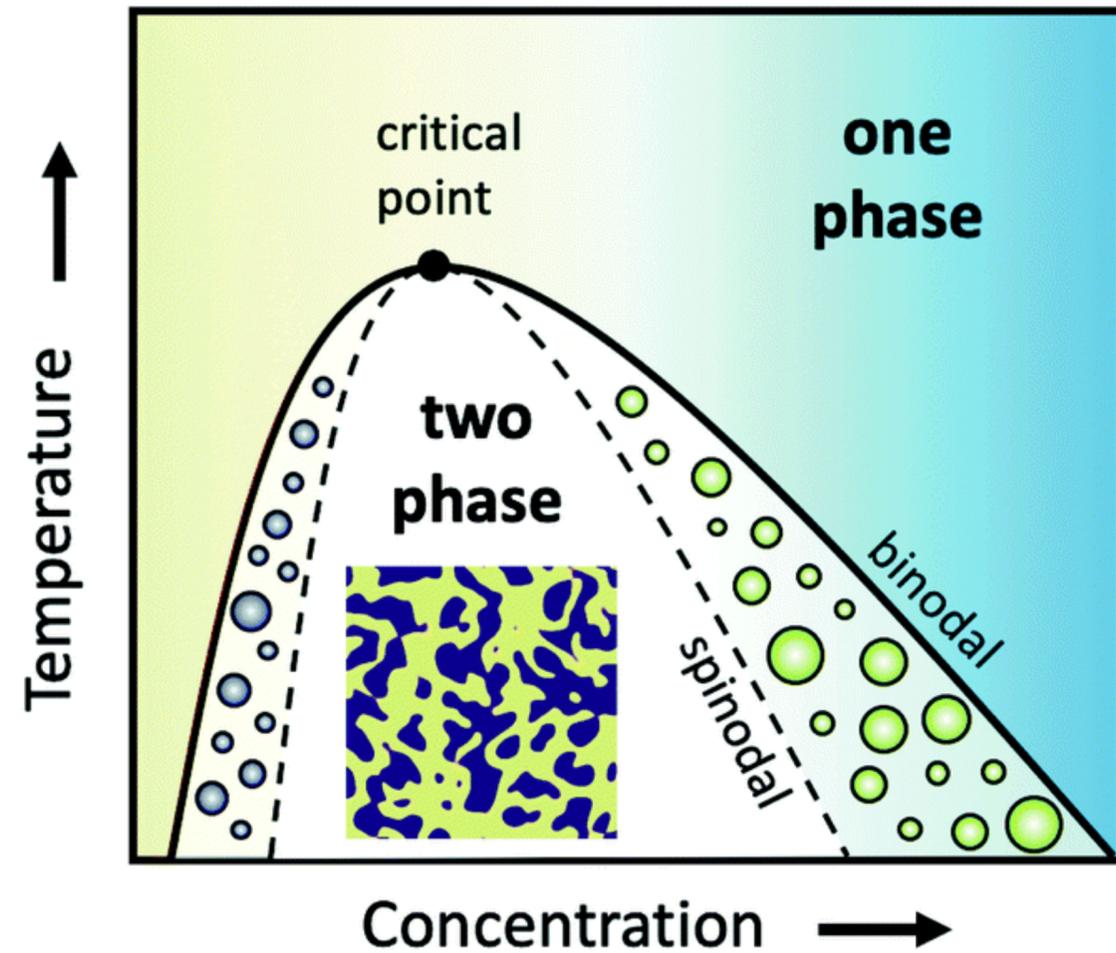
Time Scales Of Protein Diffusion



Length and time scales:
 $\tau_0(q) = 1/D_0 q^2 \Rightarrow \tau_0(1\text{nm}^{-1}) \sim 1 \mu\text{s}$

molecule	measured context	D_0 diffusion coefficient ($\mu\text{m}^2/\text{s}$)
H ₂ O	water	2000
H ₂ O	nucleus of chicken erythrocyte	200
H ⁺ (from H ₃ O ⁺ to H ₂ O)	water	7000
O ₂	water	2000
CO ₂	water	2000
tRNA (≈ 20 kDa)	water	100
protein (≈ 30 kDa GFP)	water	100
protein (≈ 30 kDa GFP)	eukaryotic cell (CHO) cytoplasm	30
protein (≈ 30 kDa GFP)	rat liver mitochondria	30
protein (NLS-EGFP)	cytoplasm of <i>D. melanogaster</i> embryo	20
protein (≈ 30 kDa)	<i>E. coli</i> cytoplasm	7-8
protein (≈ 40 kDa)	<i>E. coli</i> cytoplasm	2-4
protein ($\approx 70-250$ kDa)	<i>E. coli</i> cytoplasm	0.4-2
protein (≈ 140 kDa Tar-YFP)	<i>E. coli</i> membrane	0.2
protein (≈ 70 kDa LacY-YFP)	<i>E. coli</i> membrane	0.03
fluorescent dye (carboxy-fluorescein)	<i>A. thaliana</i> cell wall	30
fluorescent dye (carboxy-fluorescein)	<i>A. thaliana</i> mature root epidermis	3
transcription factor (LacI)	movement along DNA (1D, <i>in vitro</i>)	0.04 ($4 \times 10^5 \text{ bp}^2 \text{ s}^{-1}$)
morphogen (bicoid-GFP)	cytoplasm of <i>D. melanogaster</i> embryo	7
morphogen (wingless)	wing imaginal disk of <i>D. melanogaster</i>	0.05
mRNA	HeLa nucleus	0.03-0.10
mRNA	various localizations and sizes	0.005-1
ribosome	<i>E. coli</i>	0.04

Antibody Solutions



$$T = 25\text{ }^\circ\text{C}$$

$$\text{concentration} \approx 250\text{ mg/mL}$$

$$R_h = 5.5\text{ nm}$$

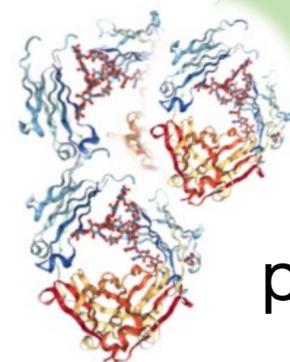
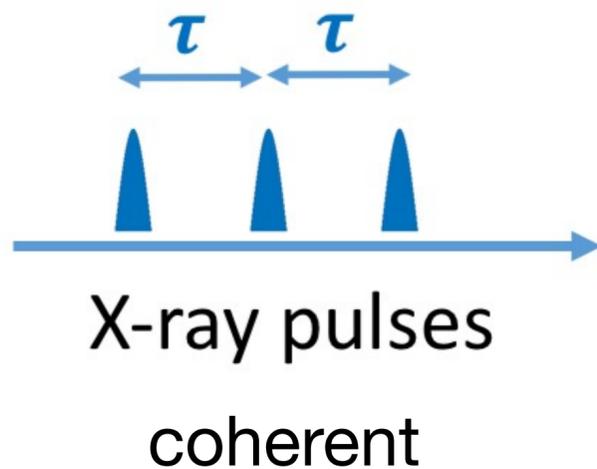
$$\tau \approx 1 - 10\text{ }\mu\text{s}$$

MHz X-Ray Photon Correlation Spectroscopy

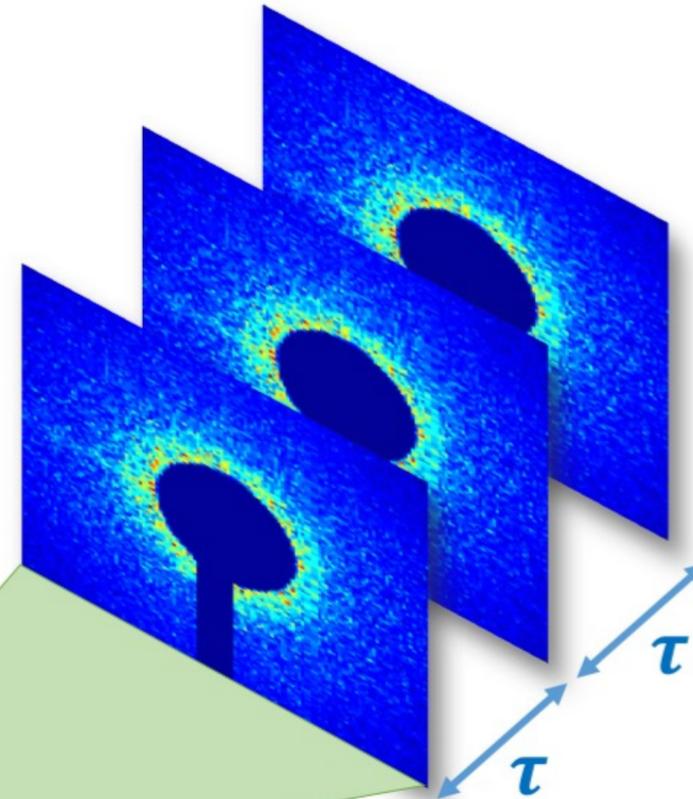
(sequential) XPCS

$$g_2(q, \tau) = \frac{\langle I(q, t)I(q, t + \tau) \rangle_t}{\langle I(q, t) \rangle_t^2}$$

Intensity auto-correlation functions



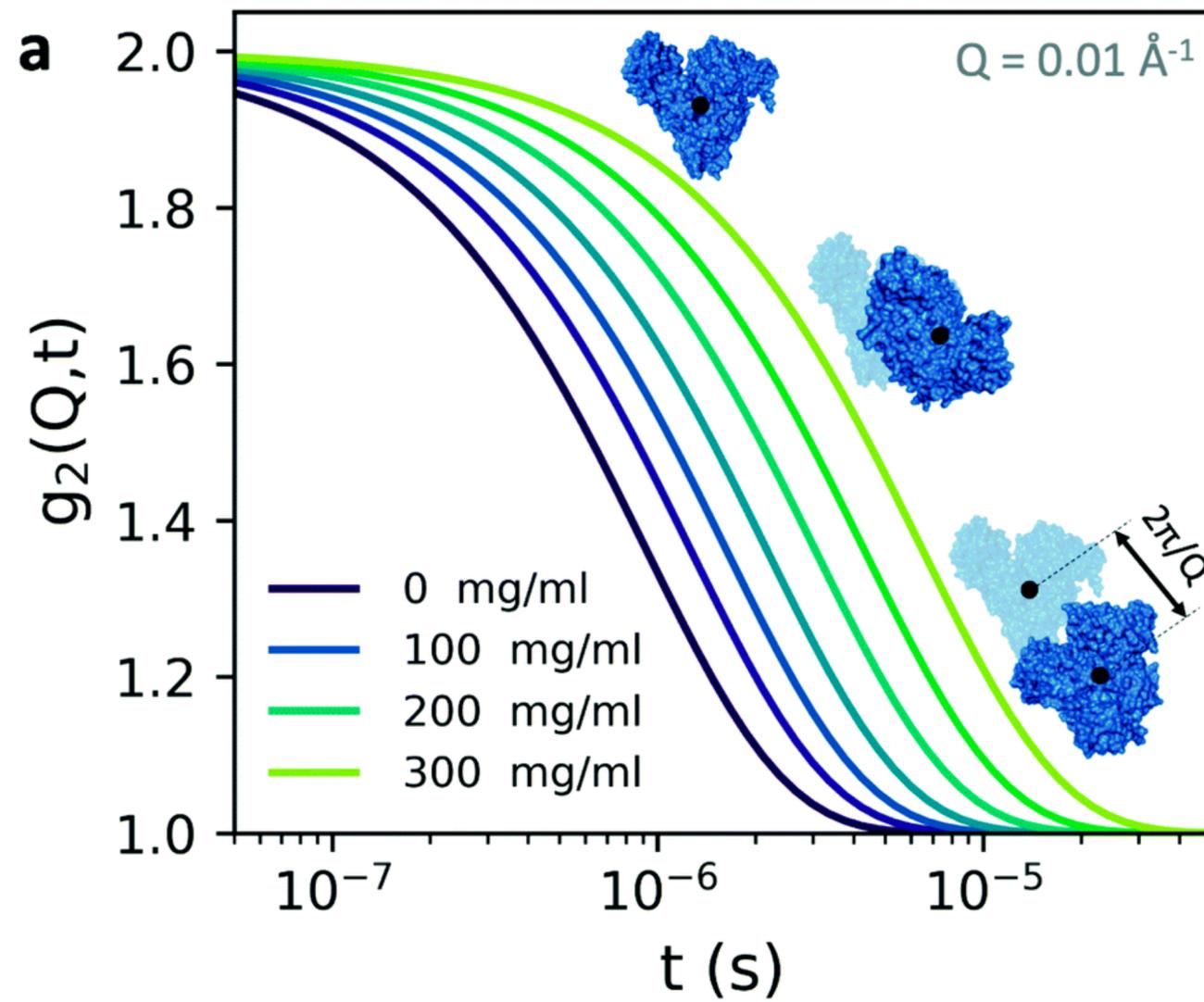
protein solution



XFEL Freq. (MHz)	temporal resolution τ (ns)
4.5	220
2.2	440
1.1	880
...	...

Measuring Protein Dynamics with MHz XPCS

Correlation Functions



$$g_2(Q, t) = 1 + e^{-2D(Q)Q^2t}$$

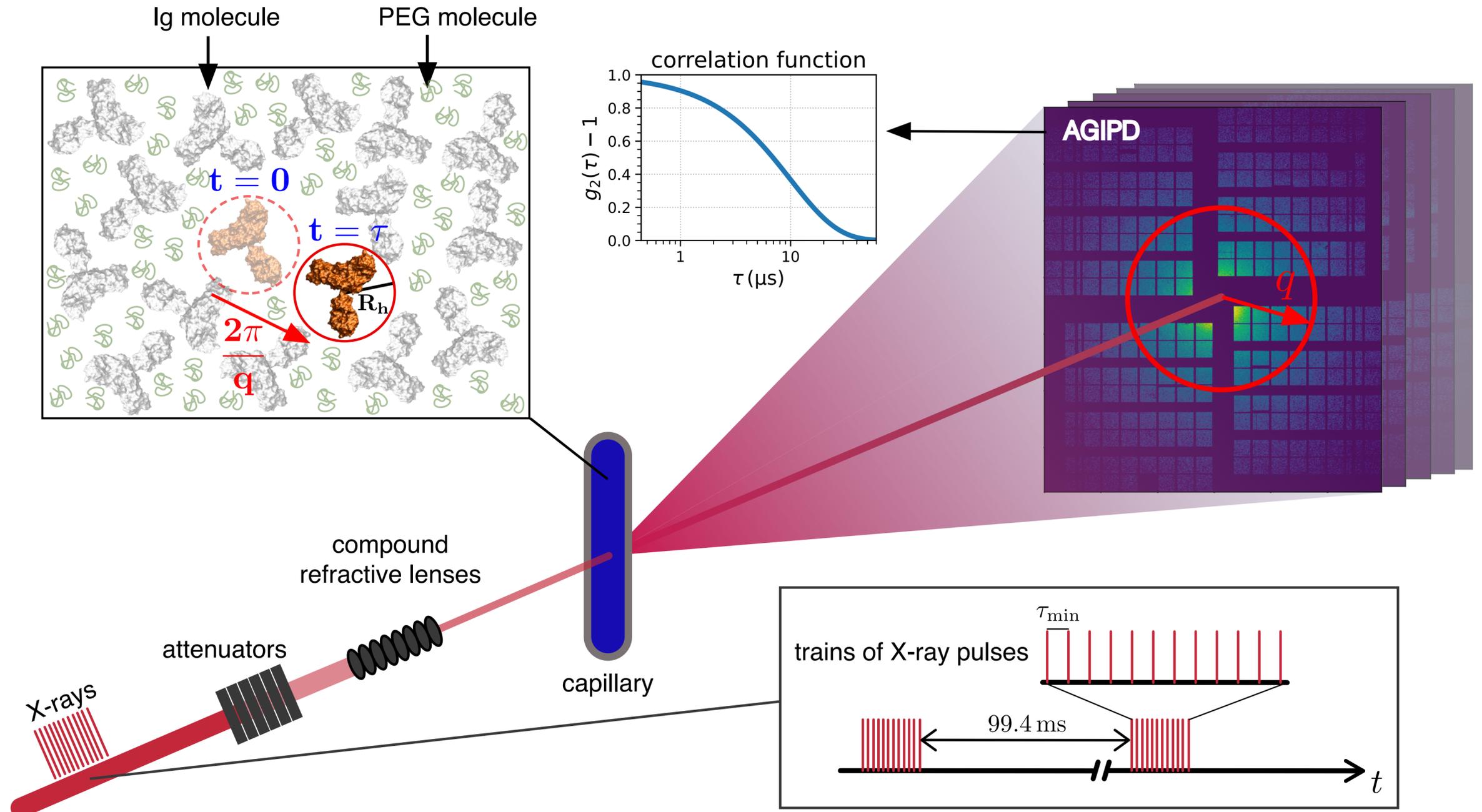
interactions

$$D(q) = \Gamma(q)/q^2 = D_0 \frac{H(q)}{S(q)}$$

SAXS MHz-XPCS Setup at MID

Setup Parameters

beam size	10 μ m x 10 μ m
attenuation	> 99 %
flux (attenuated)	1e9 photons / pulse
signal	~0.01 to 0.1 photons / pixel
sample detector distance	8m
photon energy	9keV
beam mode	pink beam
data volume	800TB
pulses per train	up to 200



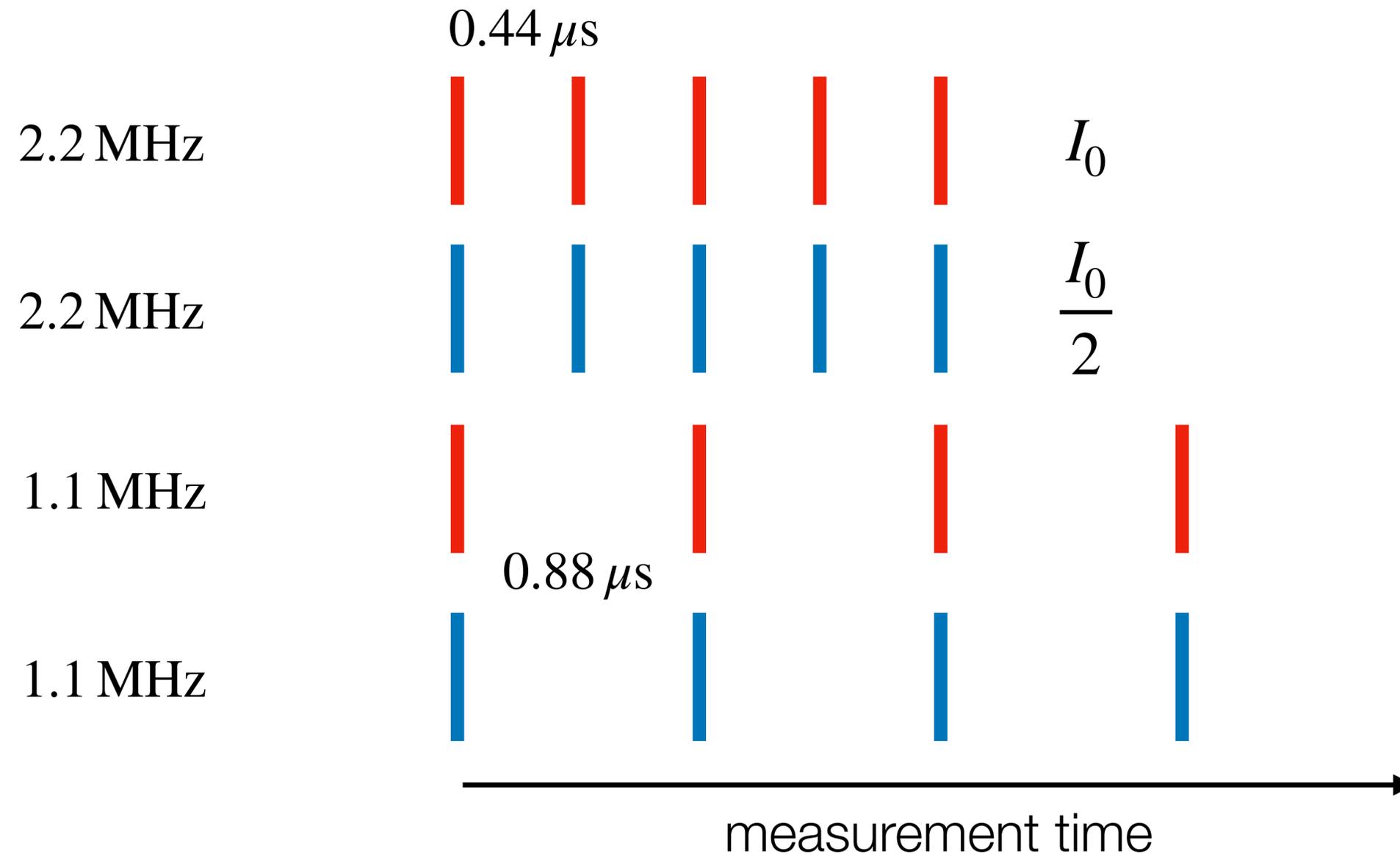
Dose and Dose Rate

MHz-XPCS with a pulsed source

Dose: 1kGy - 1kJ/kg
 Dose rate: 1kGyMHz - 1kGy/ μ s

dose
 \mathcal{D}

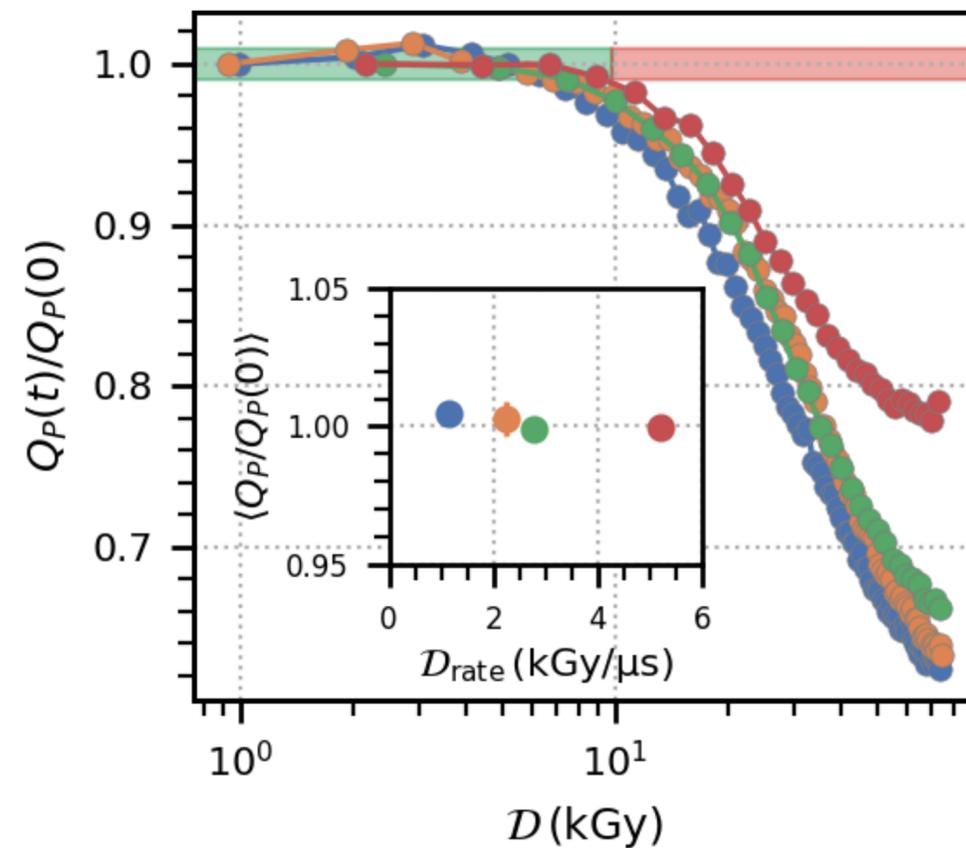
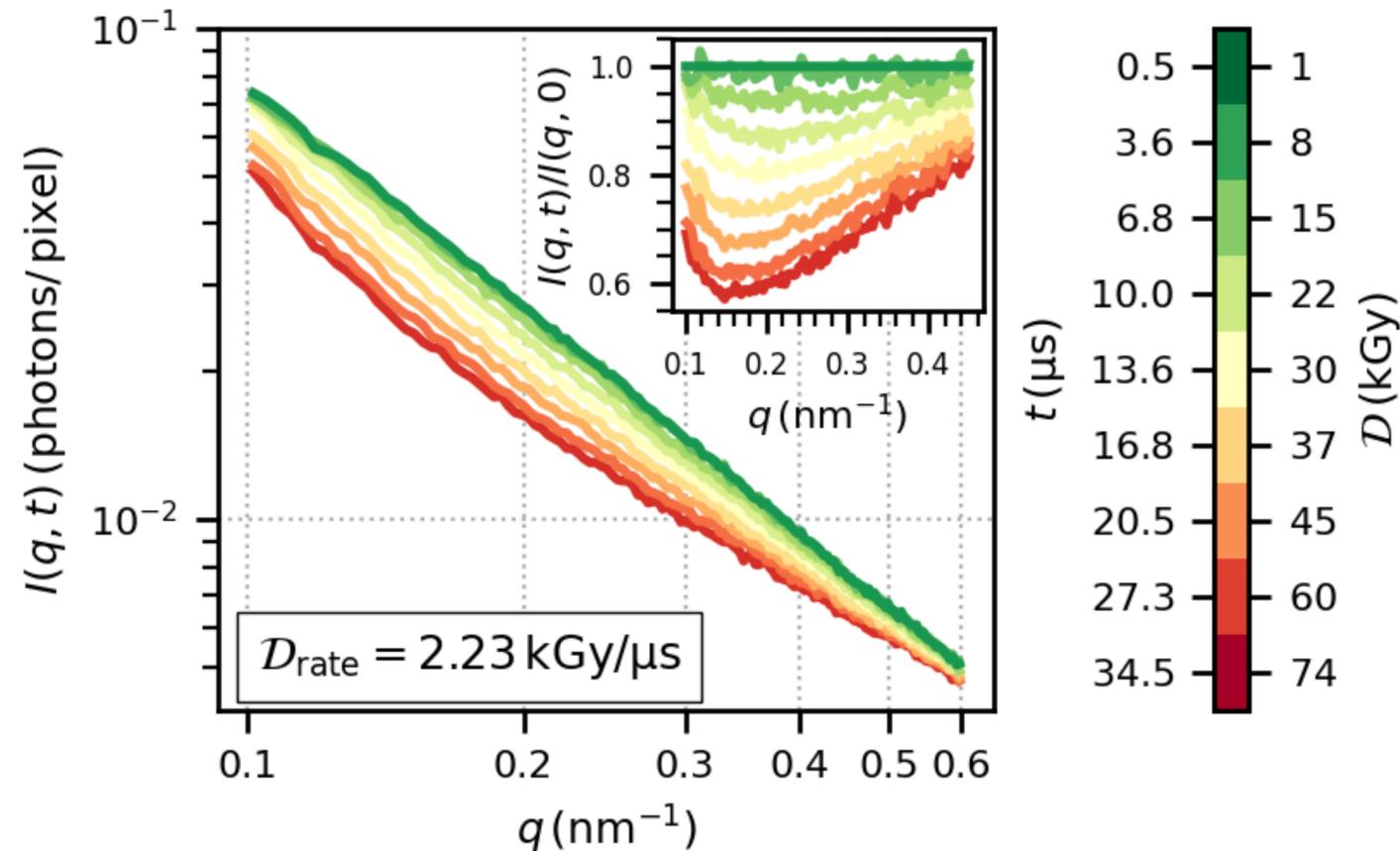
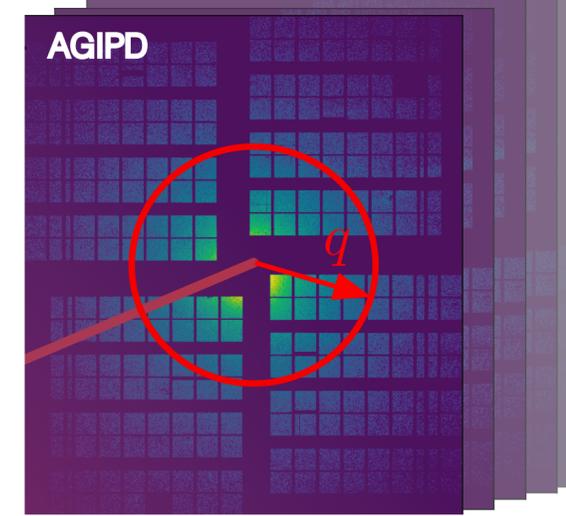
dose rate
 $\mathcal{D}_{\text{rate}}$



10 kGy	4.5 kGy/ μ s
5 kGy	2.25 kGy/ μ s
10 kGy	2.25 kGy/ μ s
5 kGy	1.23 kGy/ μ s

Structural Changes

Azimuthally Integrated Intensity



Porod invariant
$$Q_P(t) = \int_{q_{\min}}^{q_{\max}} q^2 I(q, t) dq$$

Time Resolved Dynamics

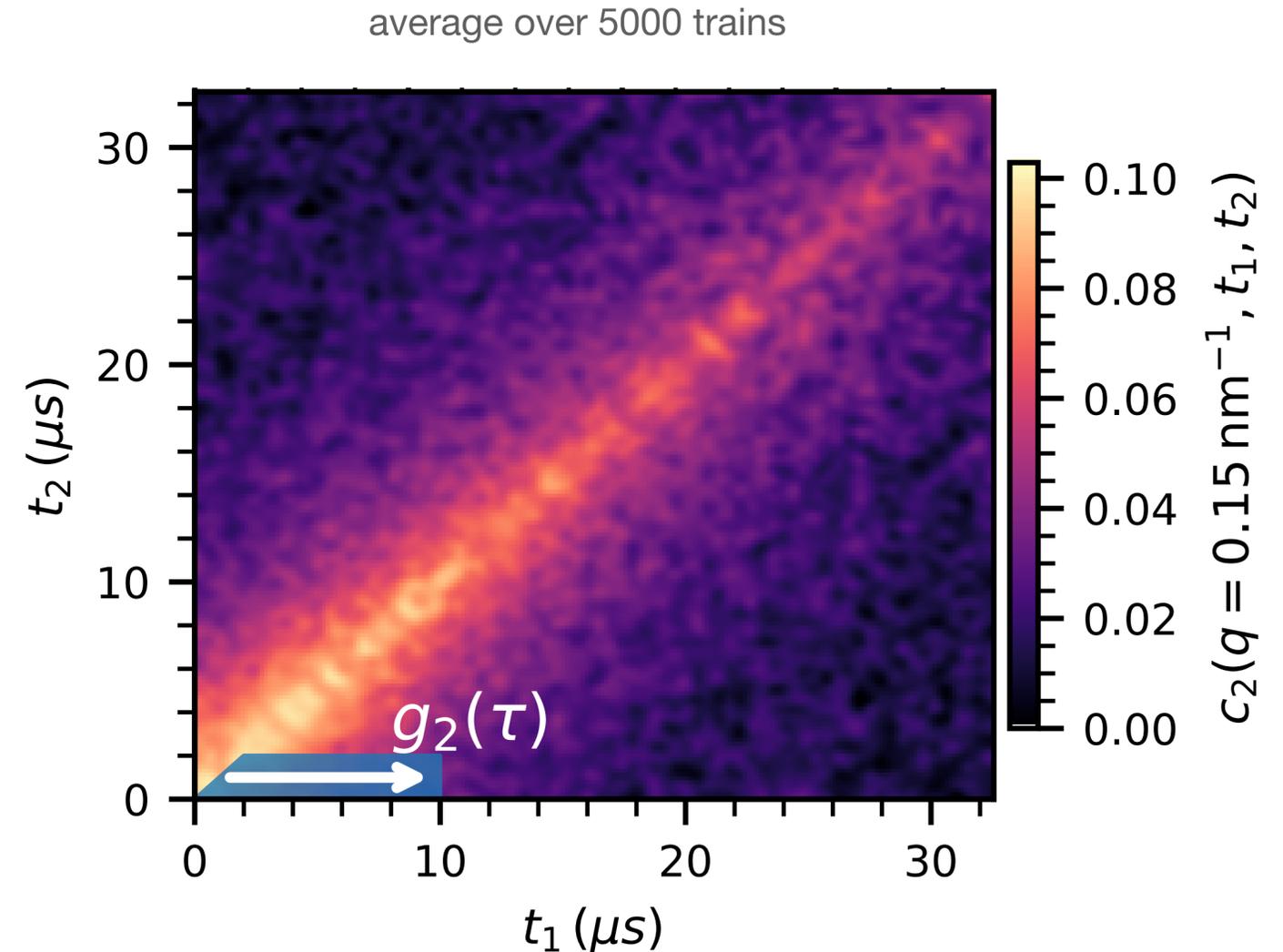
Two-Time Correlation Functions

$$g_2(q, \tau) = \frac{\langle I(q, t)I(q, t + \tau) \rangle_t}{\langle I(q, t) \rangle_t^2}$$

Intensity auto-correlation functions

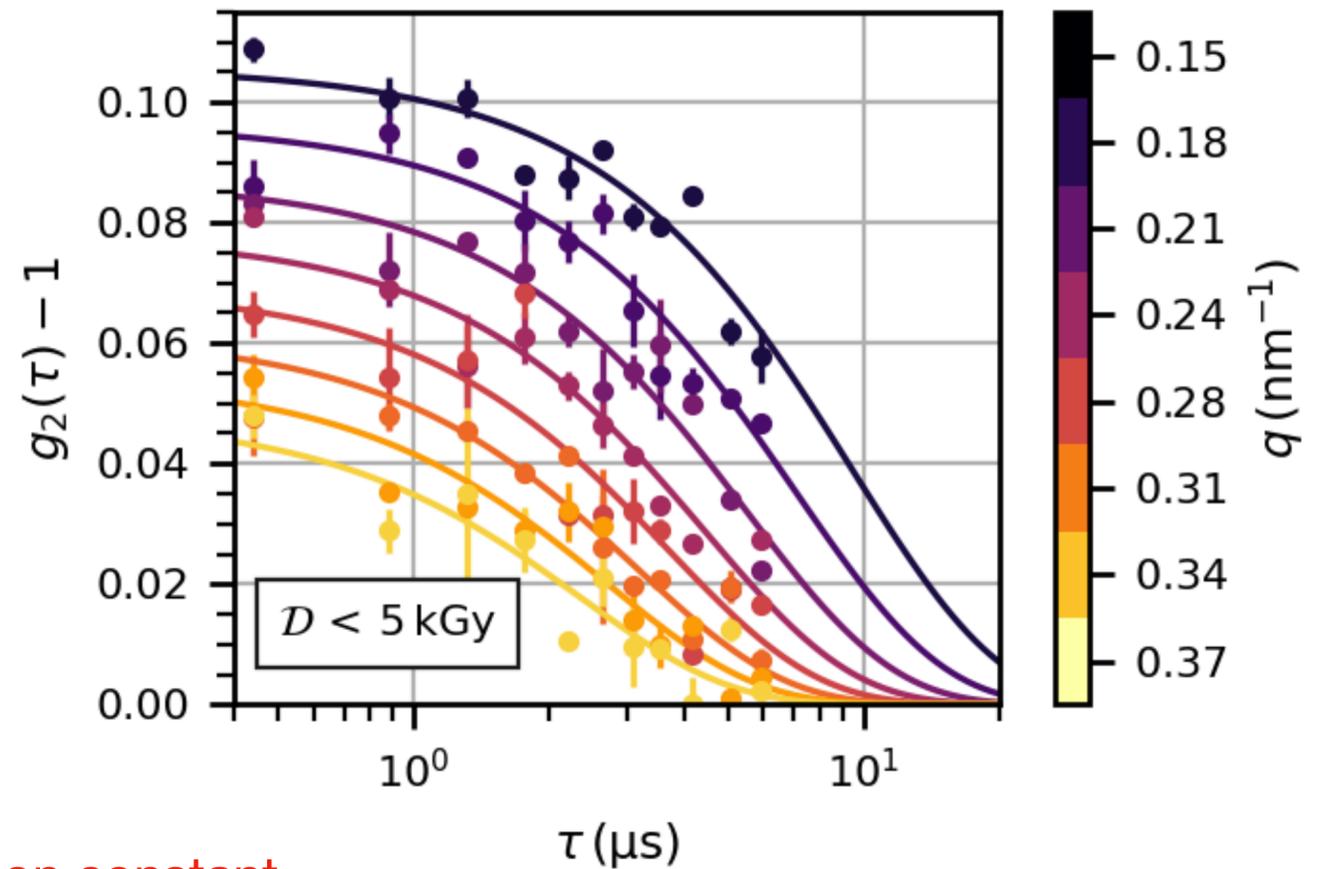
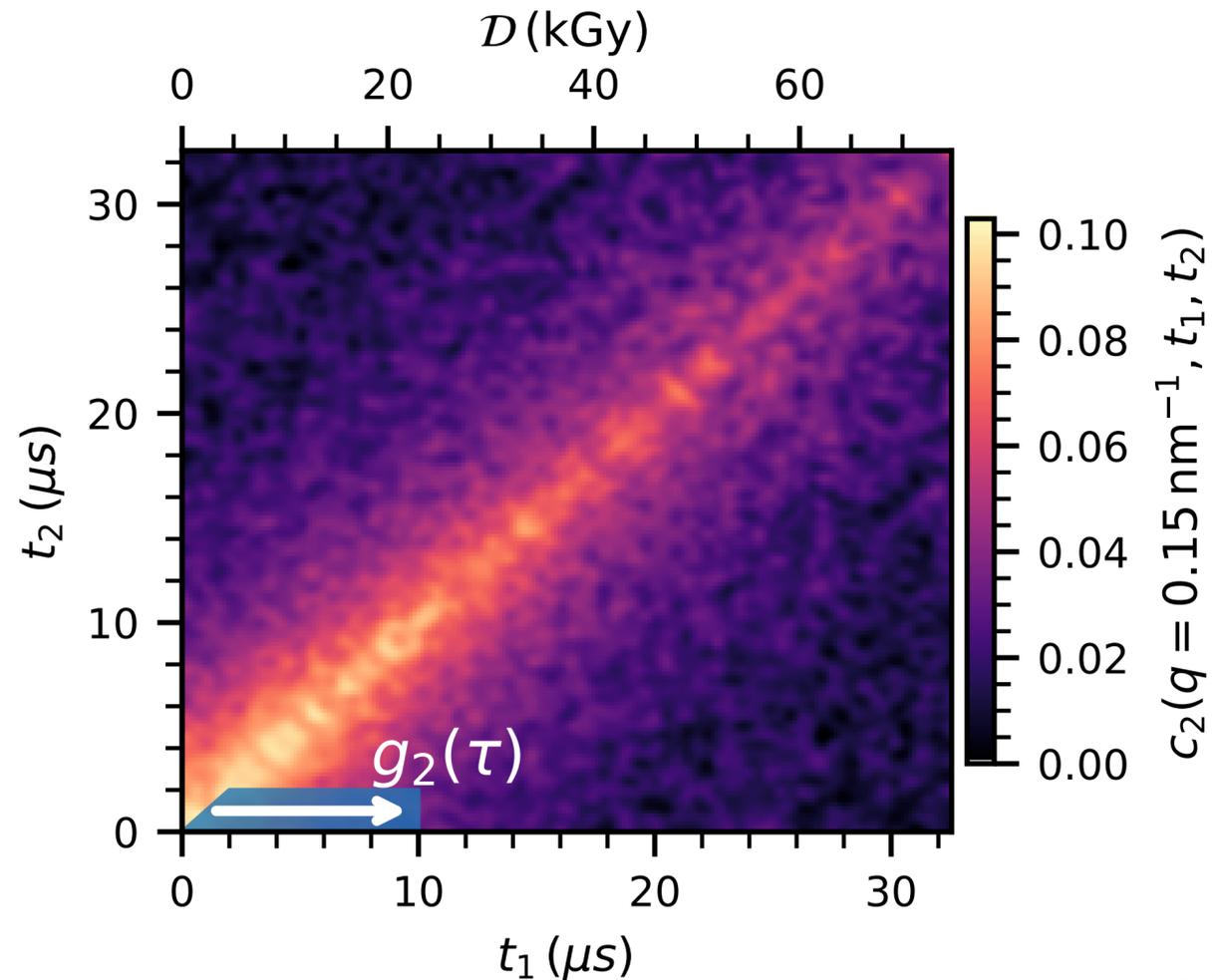
$$c_2(q, t_1, t_2) = \frac{\langle I(q, t_1)I(q, t_2) \rangle}{\langle I(q, t_1) \rangle \langle I(q, t_2) \rangle}$$

two-time correlation function



Correlation Functions

Parameter Estimation



average diffusion constant

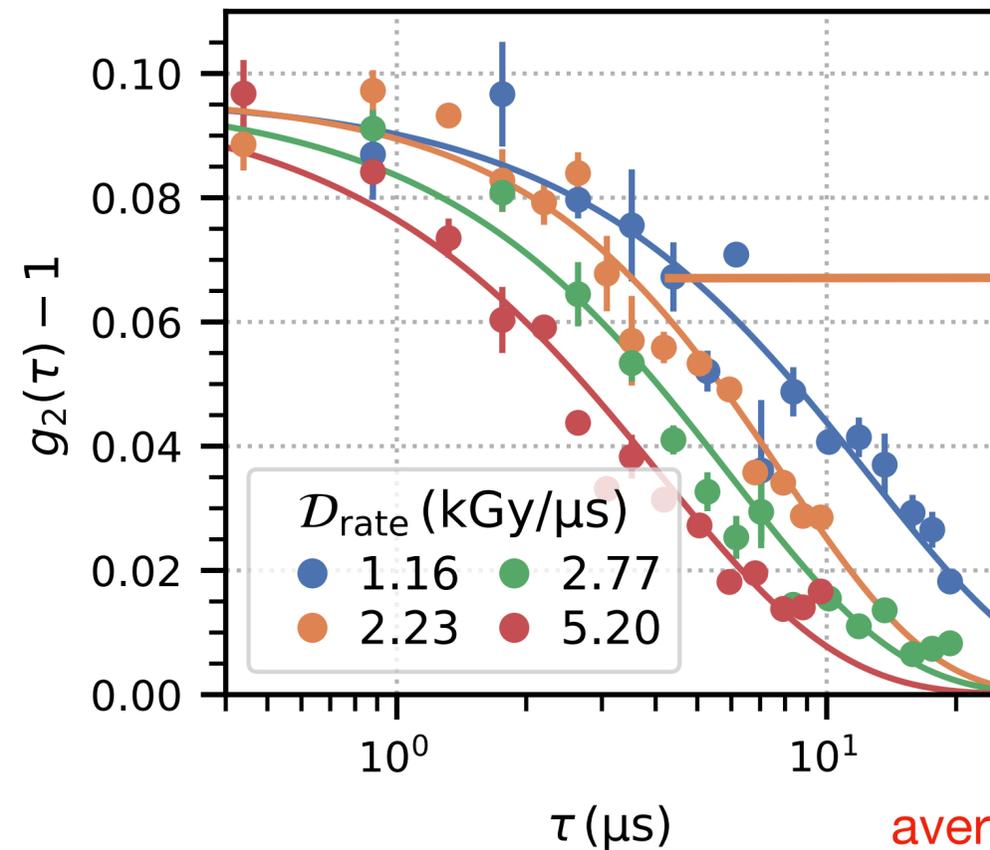
$$g_2(q, \tau) - 1 = \beta(q) e^{-2(D_0 q^2 \tau)^\alpha}$$

Kohlrausch-Williams-Watts exponent (KWW)

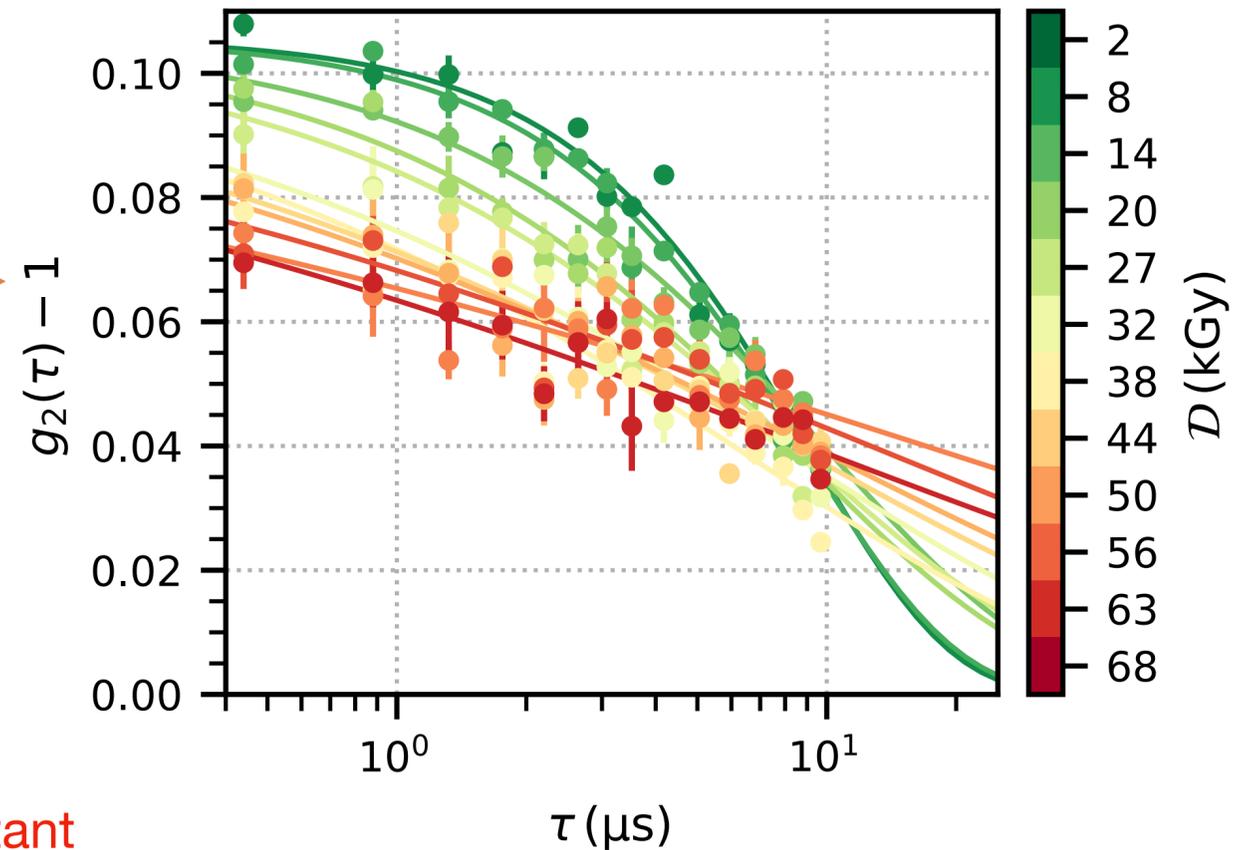
Understanding Non-Stationary Dynamics

Correlation Functions of Different Doses and Dose Rates

Dose Rate



Dose



average diffusion constant

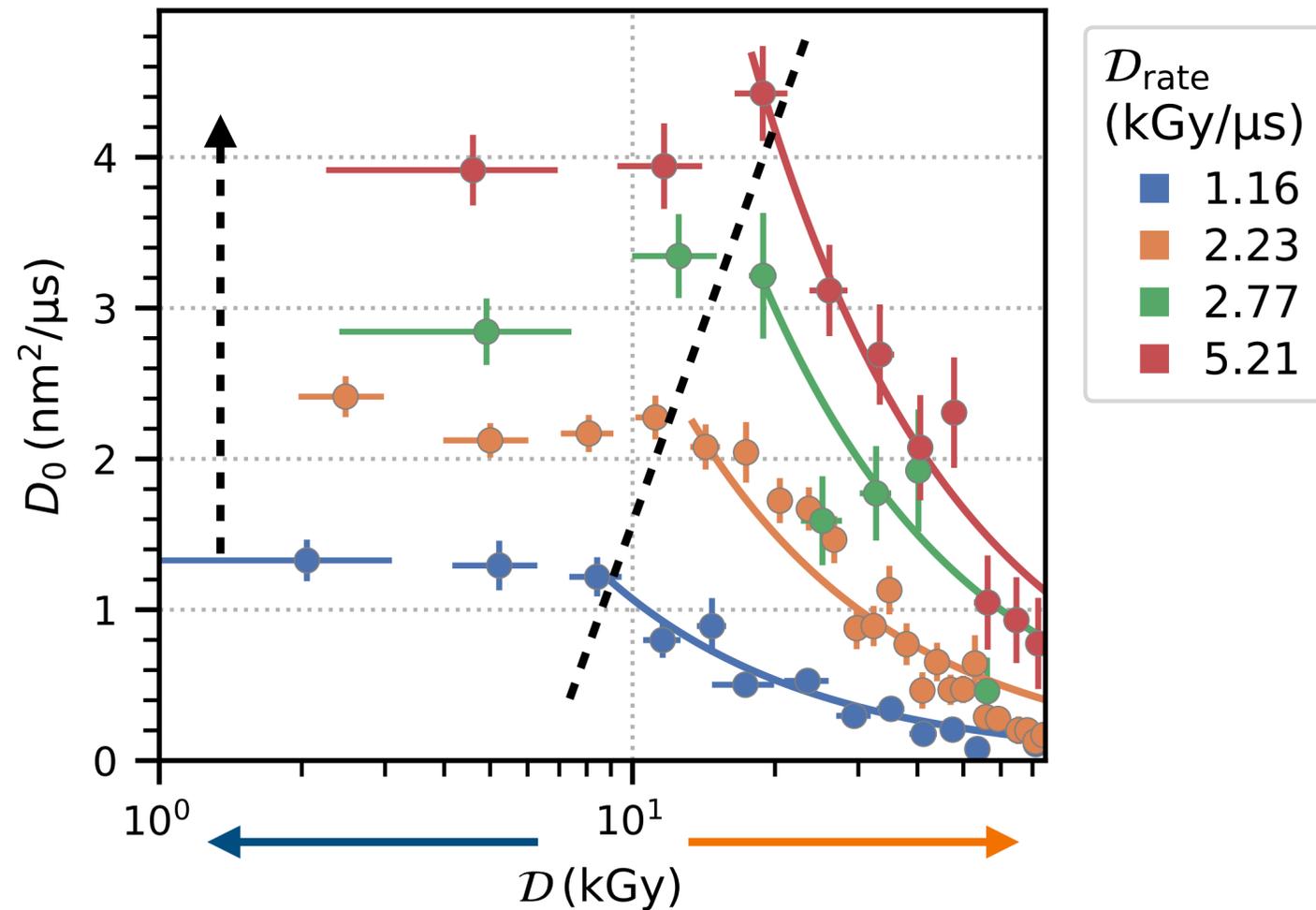
$$g_2(q, \tau) - 1 = \beta(q) e^{-2(D_0 q^2 \tau)^\alpha}$$

↙ ↘

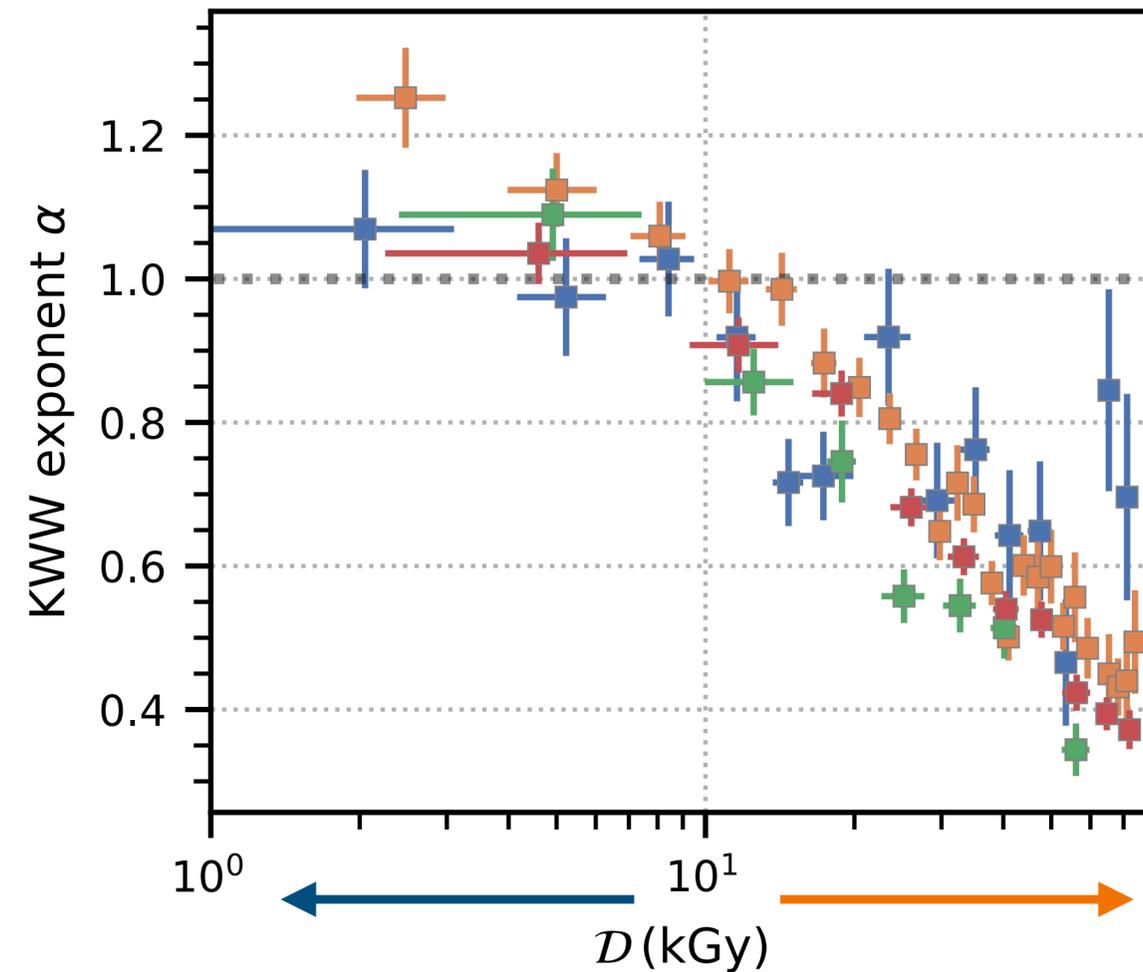
Kohlrausch-Williams-Watts exponent (KWW)

Dynamic Parameters

Diffusion Coefficient D_0



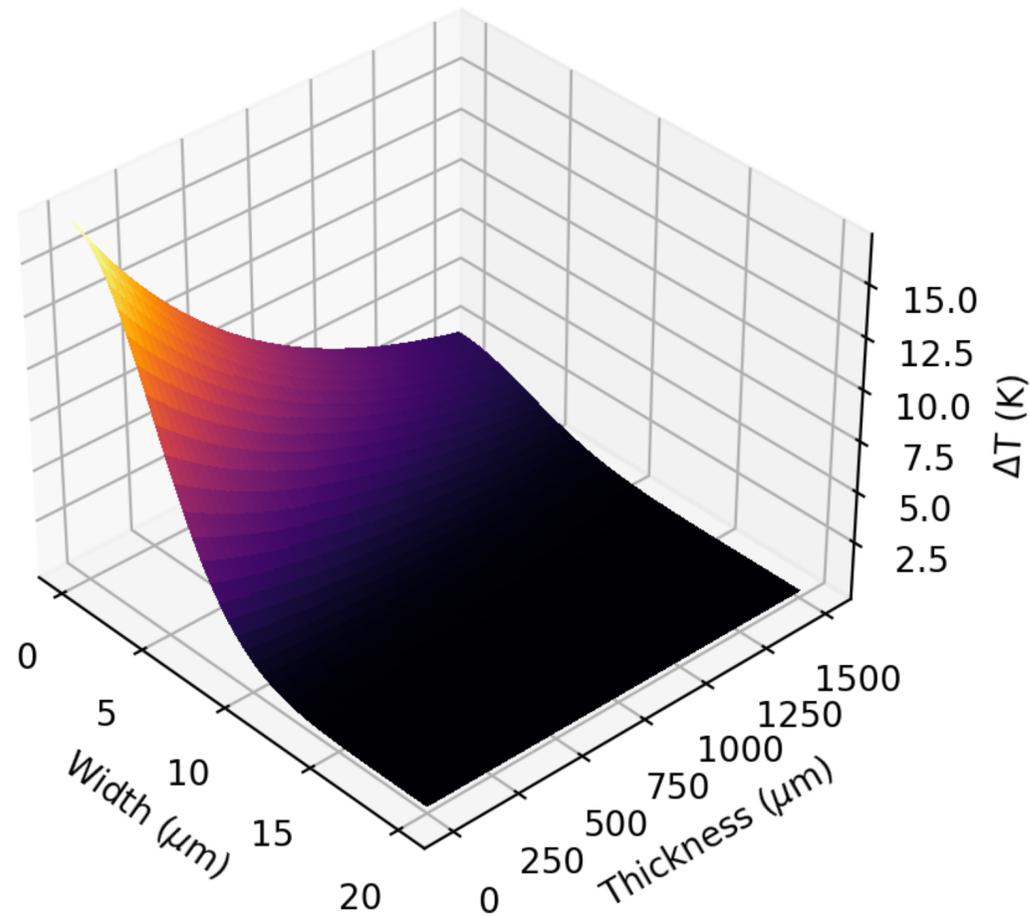
KWW exponent α



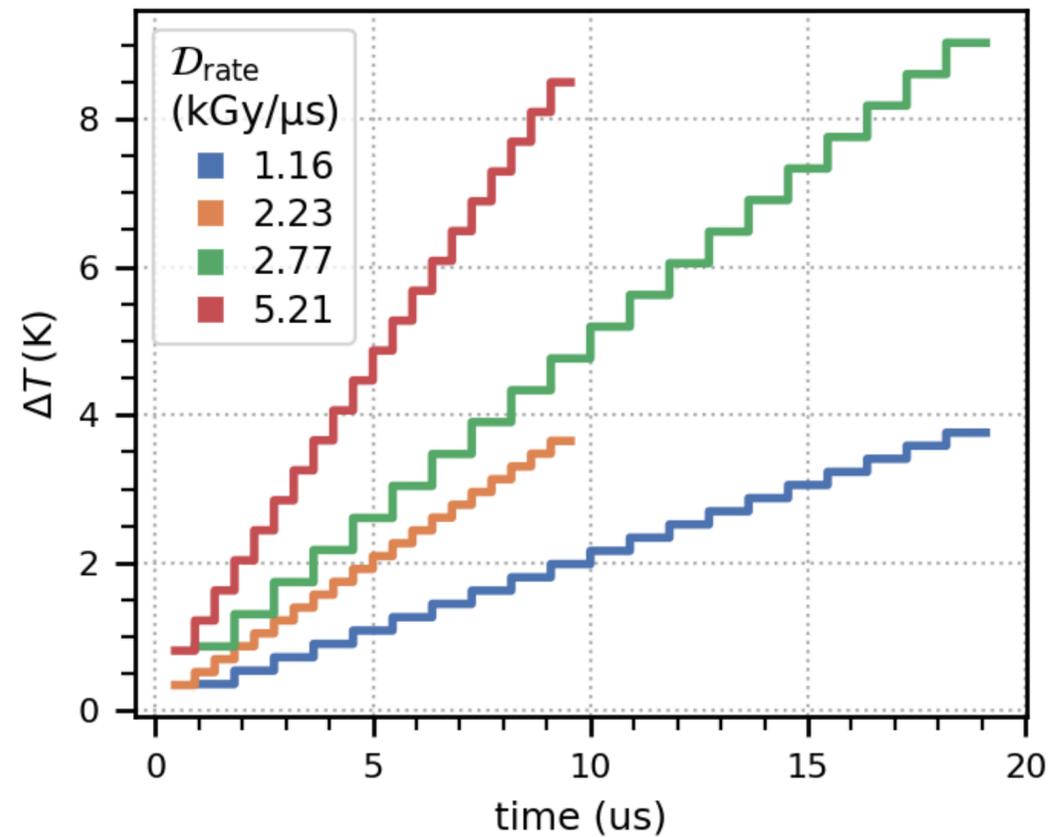
- **aggregation:** decreasing D_0 and increasing heterogeneity ($\alpha < 1$).
- **dose rate dependence:** D_0 increases with dose rate simple exponential behavior ($\alpha \approx 1$).

The Effect of X-Ray Induced Heating

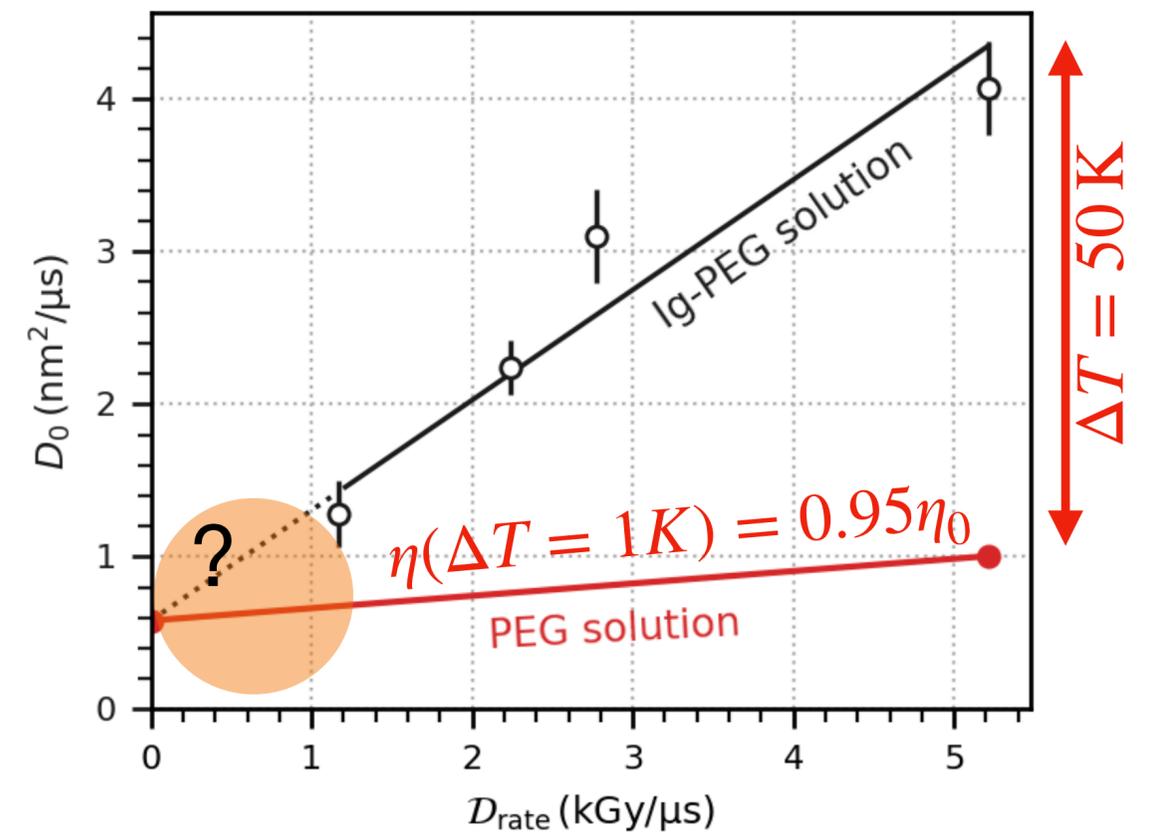
temperature profile after 20 pulses



average temperature increase

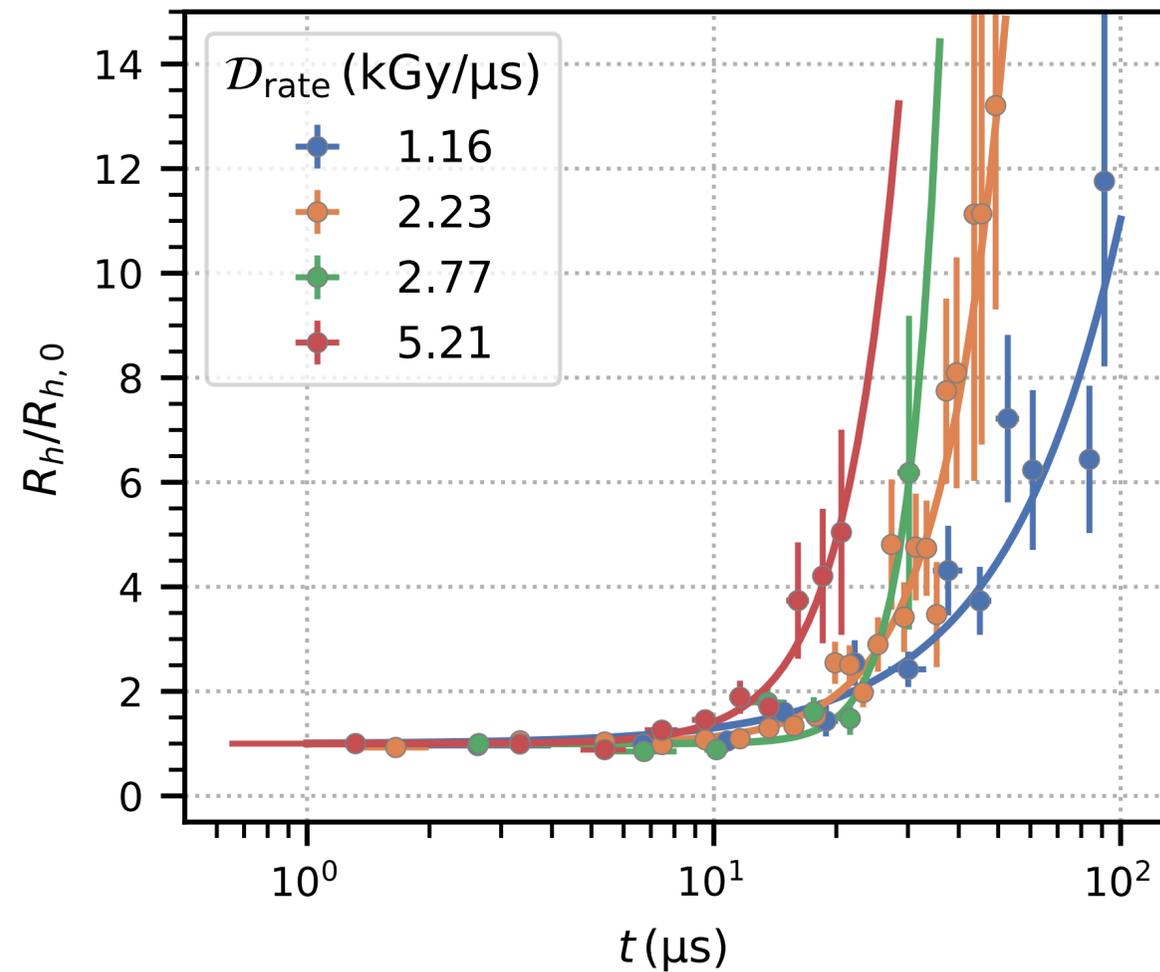


dose rate dependence of D_0

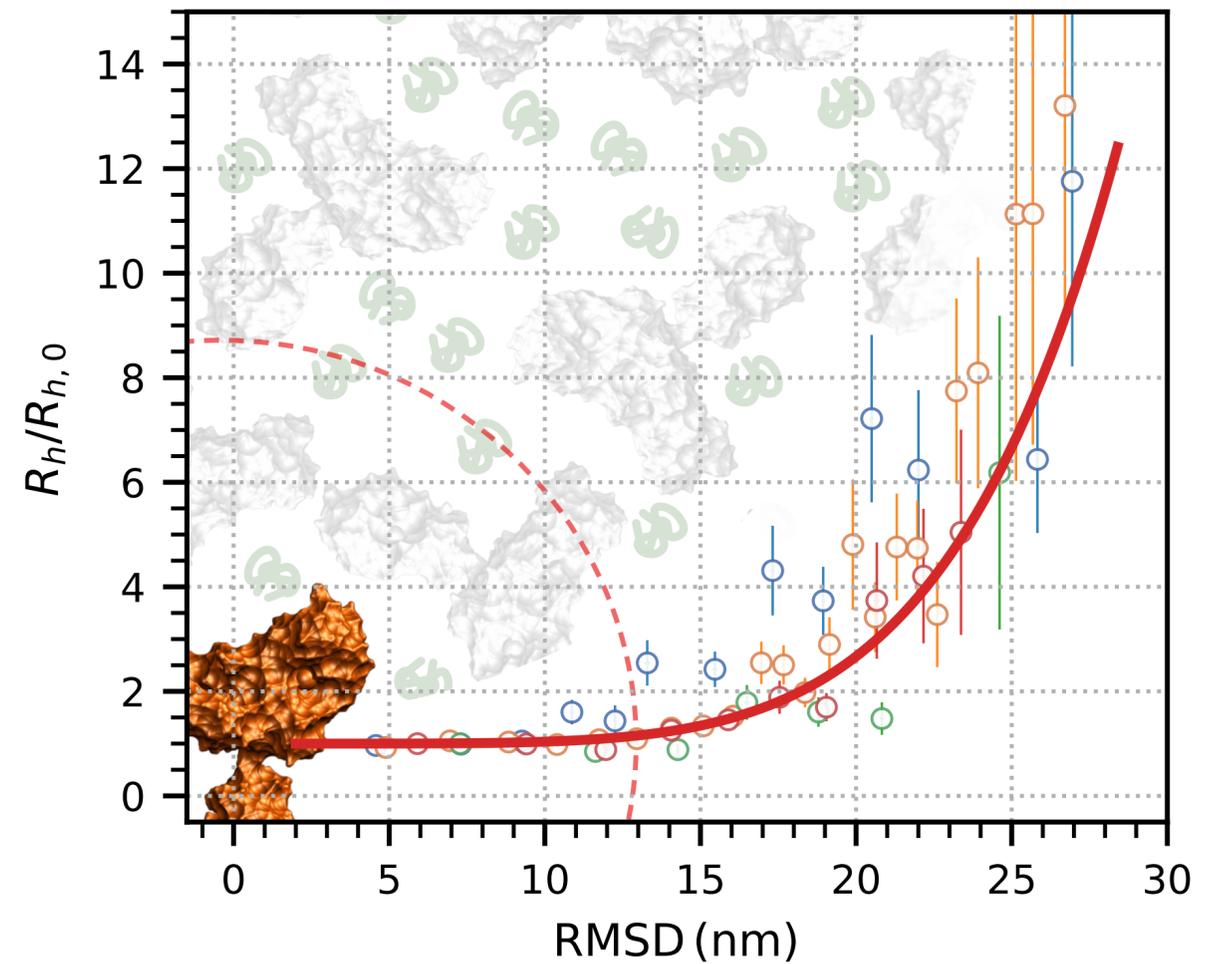


$$D_0 = \frac{k_B T}{6\pi\eta(T)R_h}$$

Correlation Before Aggregation



$$R_h = \frac{k_B T}{6\pi\eta D_0}$$



number of collisions before aggregation

$$N = \frac{\text{RMSD}^2}{\text{mean free path}^2} \approx 15$$

MHz XPCS With Antibody Solutions

First Step: The Effect of Dose and Dose Rate

- We demonstrated the feasibility of MHz-XPCS with IgG-PEG solutions.
- Distinguish between X-ray induced aggregation and speed-up.
- Dose rate dependence implies X-ray induced dynamics beyond heating.

- MHz-XPCS at XFEL would benefit from higher degree of longitudinal coherence (monochromator, seeding), faster repetition rates (4.5 MHz), longer trains (more pulses), and smaller pixels.

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Thank You.

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