

# Advancing super resolution microscopy for quantitative in-vivo imaging of chromatin nanodomains

Clayton W. Seitz

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# Outline

Introduction to fluorescence nanoscopy

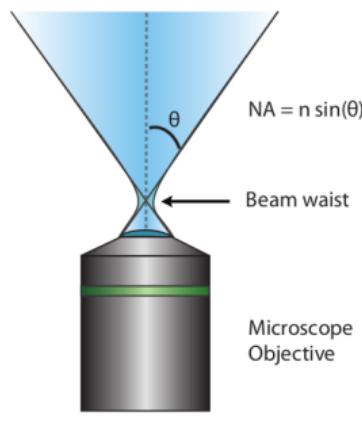
Enhanced nanoscopy with deep generative models

Super-resolution of nucleosome nanodomains *in-vivo*

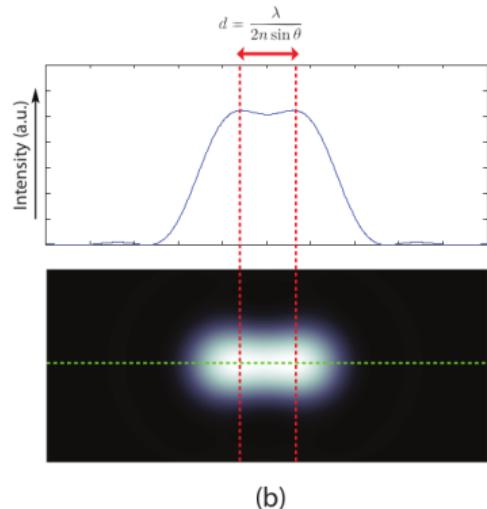
## Introduction to fluorescence nanoscopy

# Fluorescence microscopy and the diffraction limit

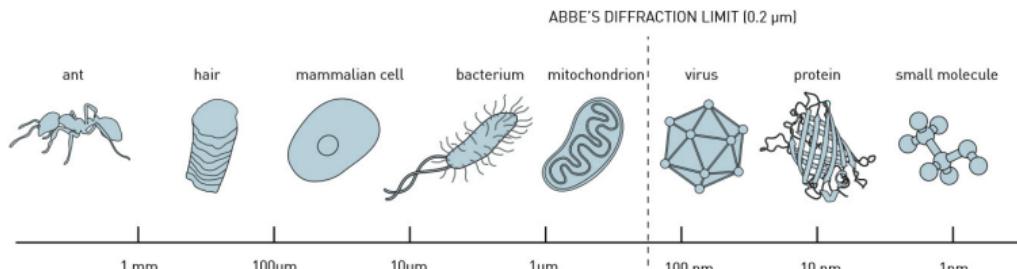
Minimal resolvable distance  $d \sim \lambda$



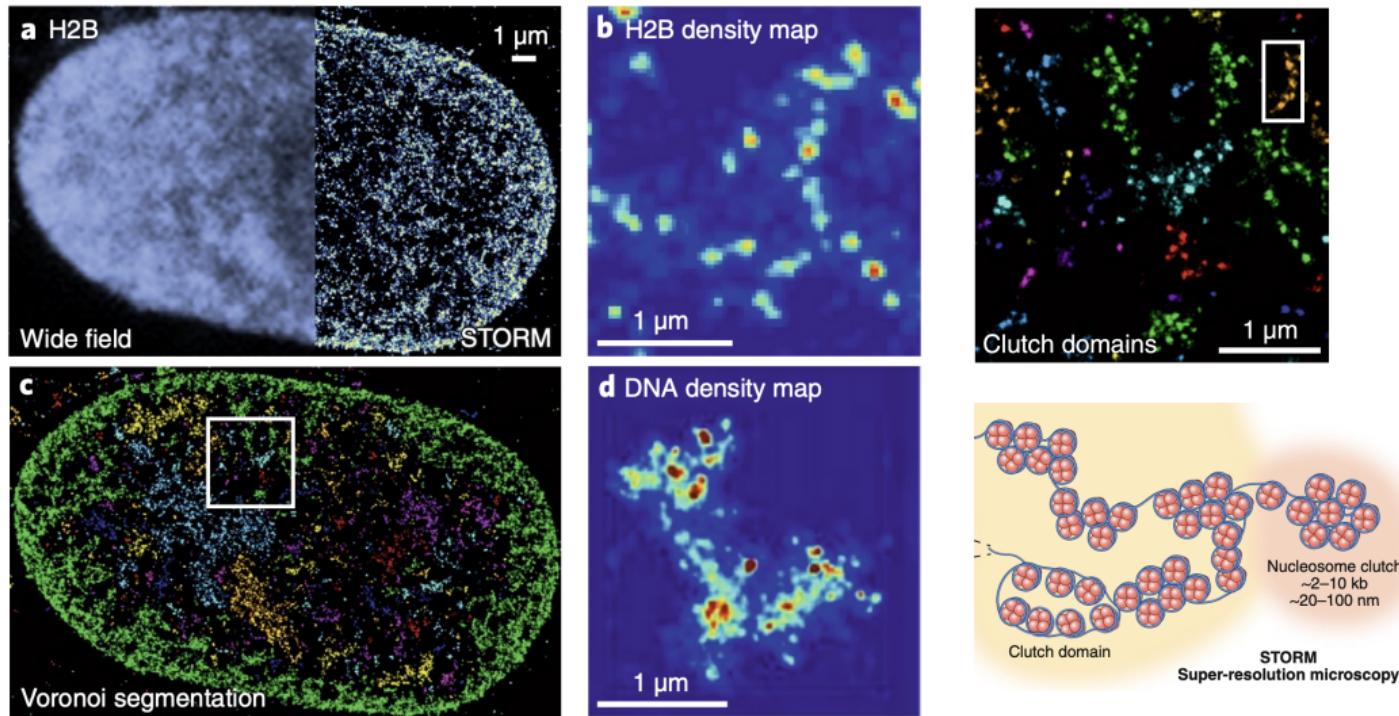
(a)



(b)



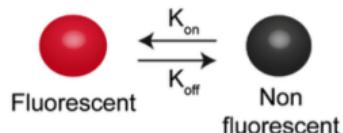
## Stochastic optical reconstruction microscopy (STORM)



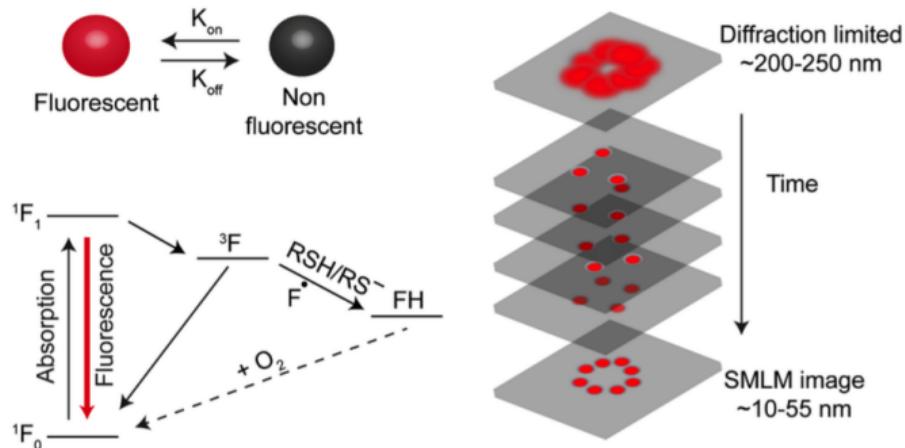
Lakadamyali, M. et al. Nature Methods 17, (2020).

# Stochastic optical reconstruction microscopy (STORM)

## a Photoswitching

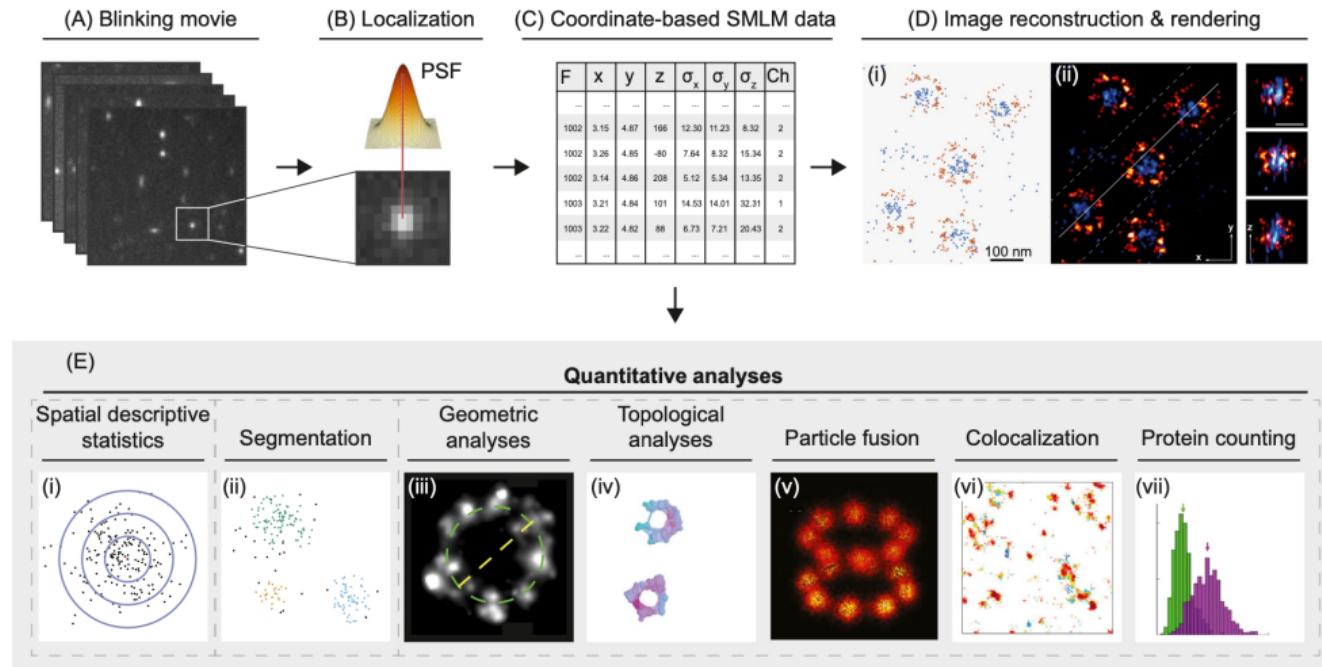


## b Temporal separation



- ▶ Single molecule localization microscopy (SMLM) techniques are diffraction-unlimited
- ▶ Photoswitching enables resolution of emitters below the diffraction limit

# Applications of single molecule localization microscopy



Wu et al. Trends in Cell Biology. 30 (2020)

## Enhanced nanoscopy with deep generative models

# Localization of isolated fluorescent emitters

Modeling the point spread function permits sub-pixel localization

$$\mu_k = i_0 \int \int O(u, v) du dv$$

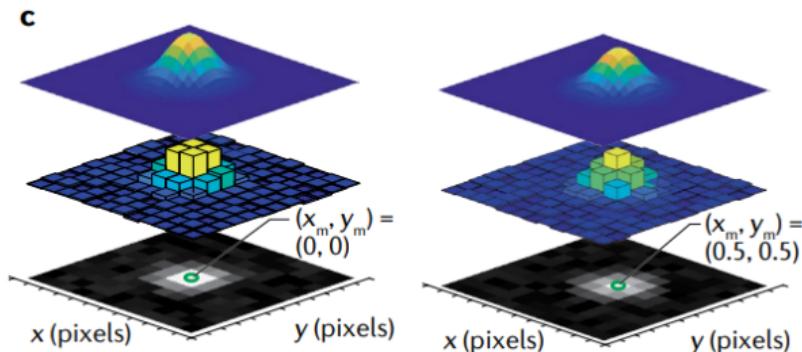
$$i_0 = g_k \eta N_0 \Delta$$

$g_k$  – pixel gain

$\eta$  – quantum efficiency

$N_0$  – photon emission rate

$\Delta$  – exposure time

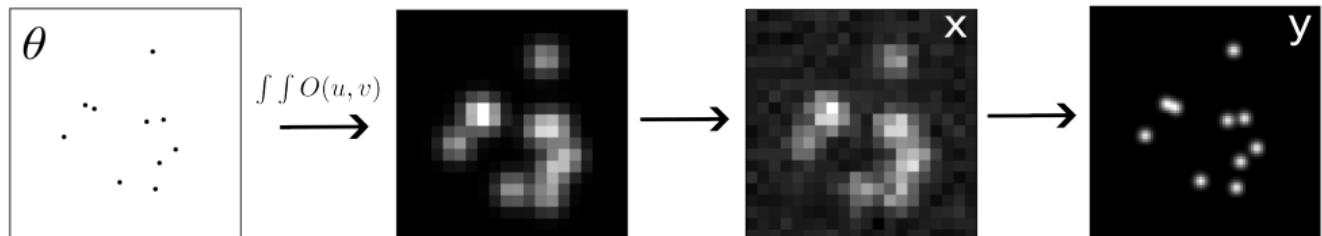


Assume  $N_0$  is constant over  $\Delta$  (homogeneous Poisson)

$$\theta^* = \operatorname{argmax}_{\theta} \prod_k P(\mathbf{x}_k | \theta) = \operatorname{argmin}_{\theta} - \sum_k \log P(\mathbf{x}_k | \theta)$$

# How to pack more localizations in a single frame?

Cast localization as *image restoration*



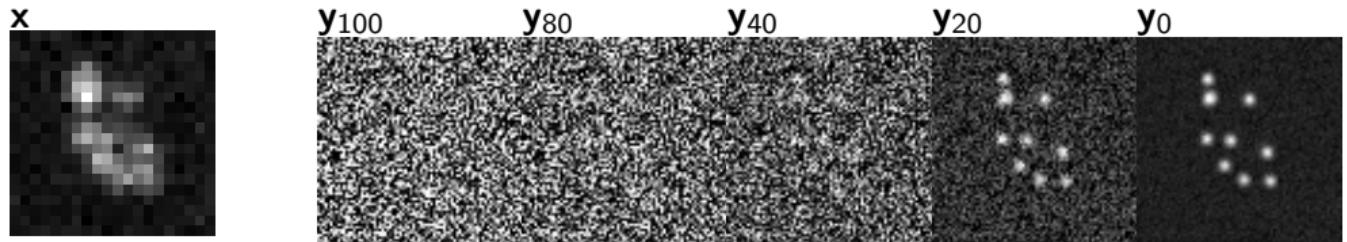
Seitz et al. Under Review. (2024)

- ▶ Would like to estimate a high-resolution  $\mathbf{y}$  from low-resolution  $\mathbf{x}$ , but it is many to one
- ▶ Must then model a *distribution* over  $\mathbf{y}$  i.e.,  $p(\mathbf{y}|\mathbf{x})$
- ▶ How to model a distribution over images?

## Bayesian image restoration with diffusion models

$$\mathbf{y}_{t-1} = \frac{1}{\sqrt{1 - \beta_t}} (\mathbf{y}_t + \beta_t s_\theta(\mathbf{y}_t)) + \sqrt{\beta_t} \xi \quad \xi \sim \mathcal{N}(0, I)$$

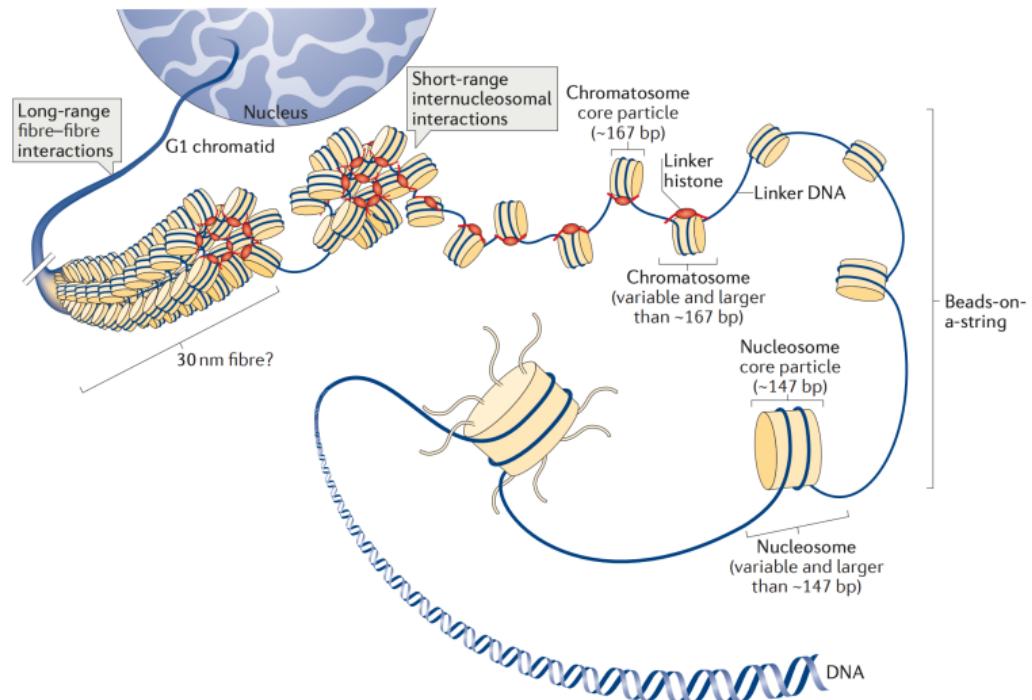
# Bayesian image restoration with diffusion models



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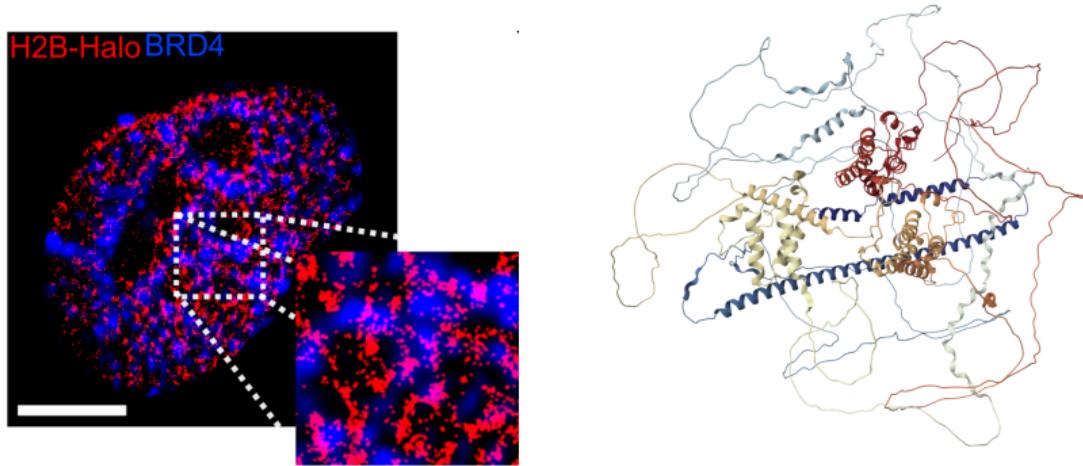
## Super-resolution of nucleosome nanodomains *in-vivo*

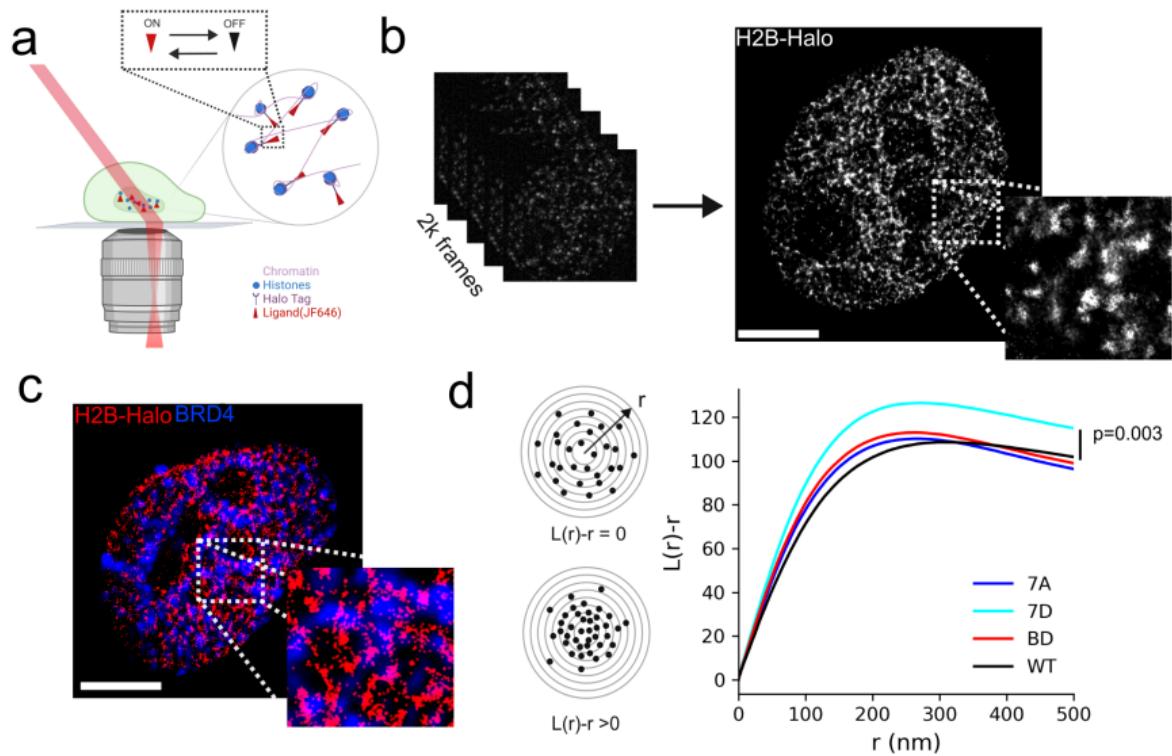
# Hierarchical structure of chromatin



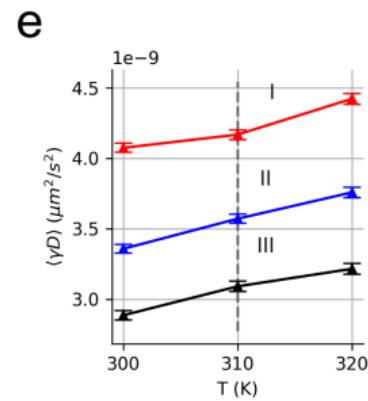
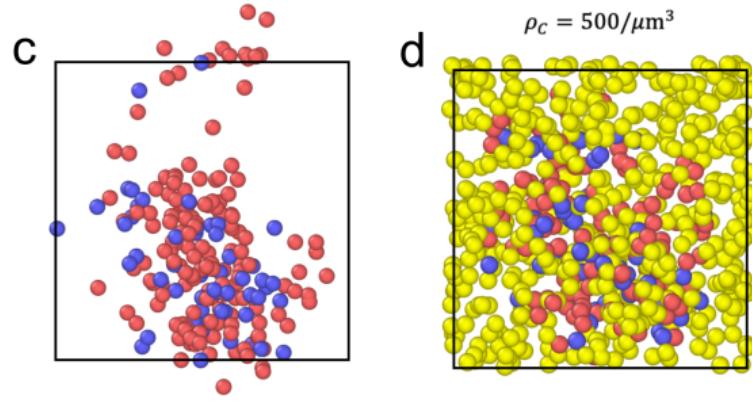
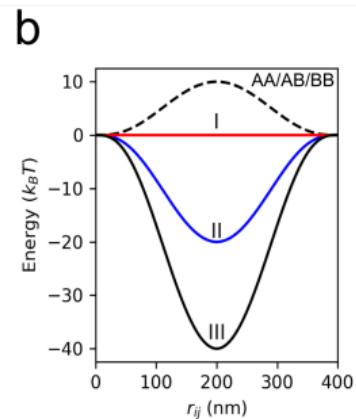
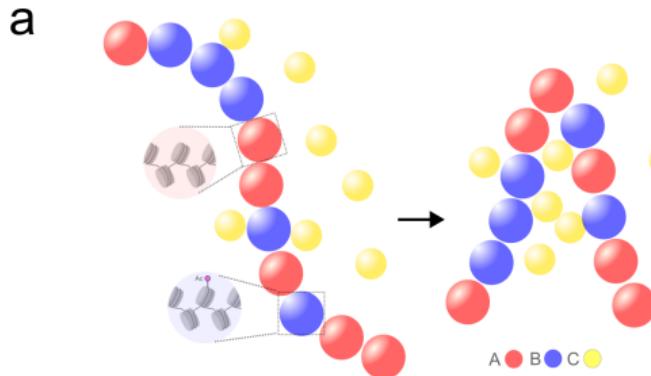
Fyodorov, D. et al. Nat Rev Mol Cell Biol 19, (2018).

# Bromodomain protein 4 (BRD4) binds acetylated chromatin

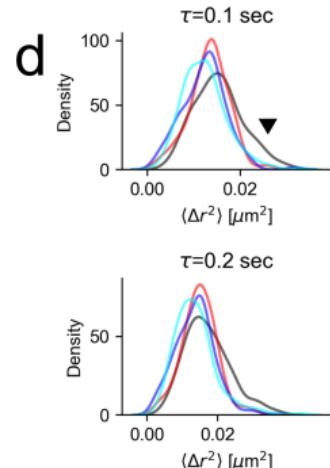
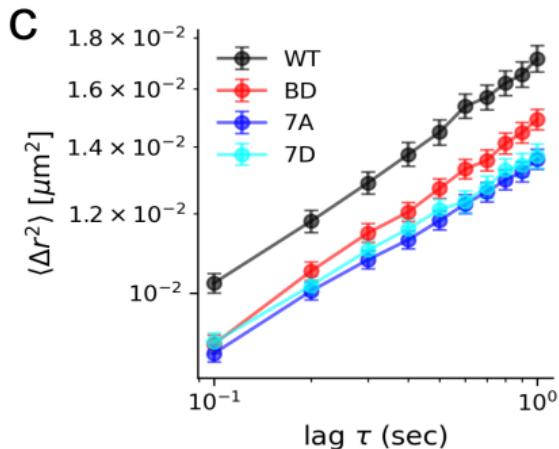
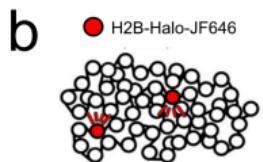
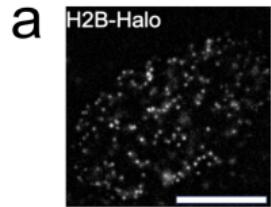




Seitz et al. Under Review. (2020)



Seitz et al. Under Review. (2020)



Seitz et al. Under Review. (2024)

## Recent Publications

- ▶ Maelle Locatelli<sup>†</sup>, Josh Lawrimore<sup>†</sup>, Hua Lin<sup>†</sup>, Sarvath Sanaullah, **Clayton Seitz**, ..., Pierre-Alexandre Vidi. *DNA damage reduces heterogeneity and coherence of chromatin motions.* PNAS. July 2022
- ▶ Mengdi Zhang, **Clayton Seitz**, Garrick Chang, Fadil Iqbal, Hua Lin, and Jing Liu *A guide for single-particle chromatin tracking in live cell nuclei.* Cell Biology International. January 2022.
- ▶ Wenting Wu, Farooq Syed, Edward Simpson, Chih-Chun Lee, Jing Liu, Garrick Chang, Chuanpeng Dong, **Clayton Seitz**, ..., Carmella Evans-Molina; *Impact of Proinflammatory Cytokines on Alternative Splicing Patterns in Human Islets.* Diabetes. January 2022

# Acknowledgements



(left to right) Charles Park, Garrick Chang, Jing Liu, David Buchanan, Mengyuan Liu, Hailan Ma



Pancho



Donghong Fu



Norbert Scherer

Thank you!