

Sriram Fest



Happy Anniversary!



Examples of Fluctuations and microphase separation in Nuclei

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The characteristics of nuclear membrane fluctuations in stem cells

Sedigheh Ghanbarzadeh Nodehi^a, G. V. Shivashankar^{b,c}, Jacques Prost^{d,e,1}, and Farshid Mohammad-Rafiee^{a,f,1}

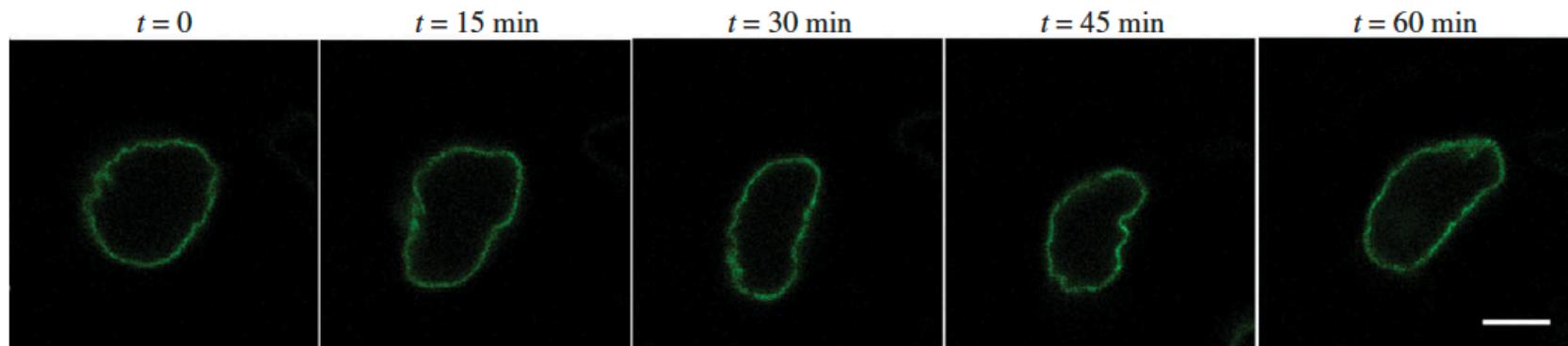
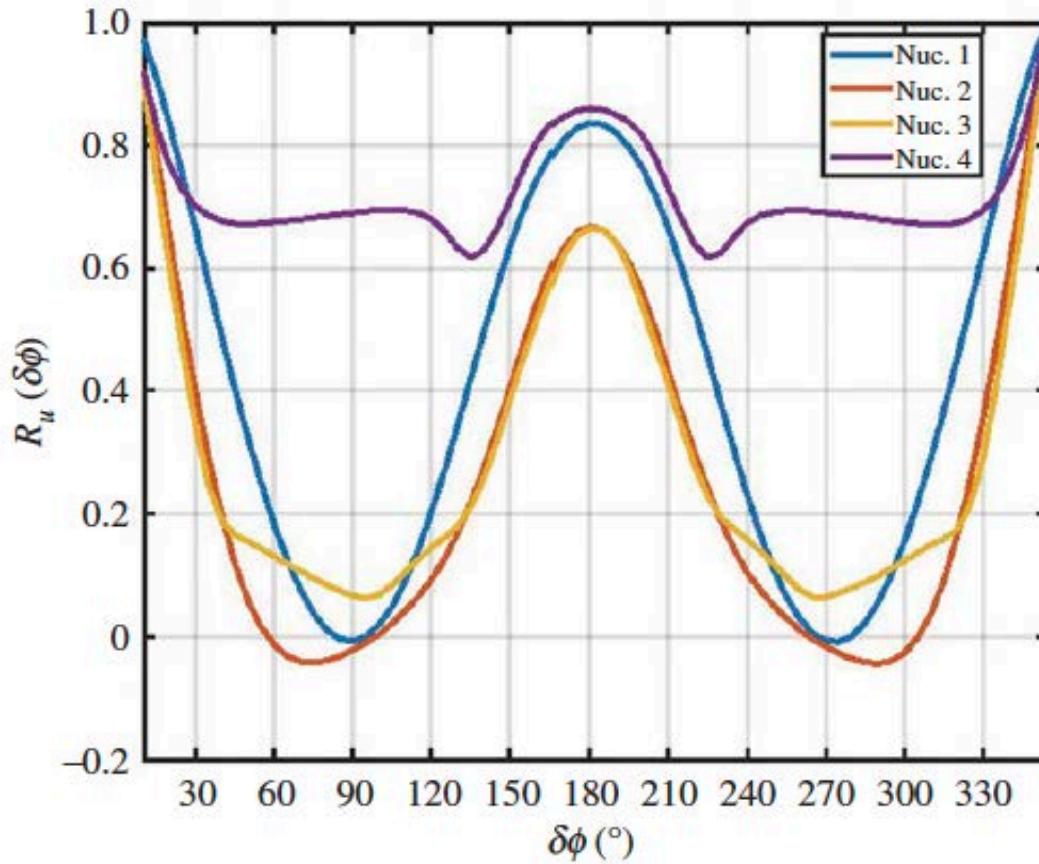


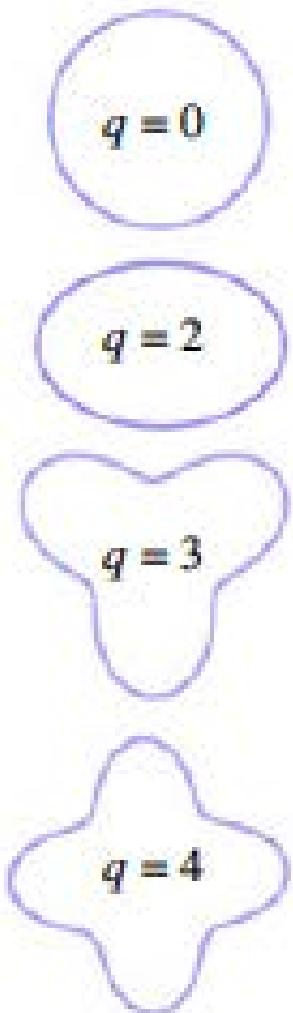
Figure 1. Stained membrane of the nucleus of an ES cell [16] over time with 15 min time intervals is shown. The scale bar is 5 μm . The total image sequences are in 10 s time resolution and reveal the highly dynamic nature of the ES cell nucleus.



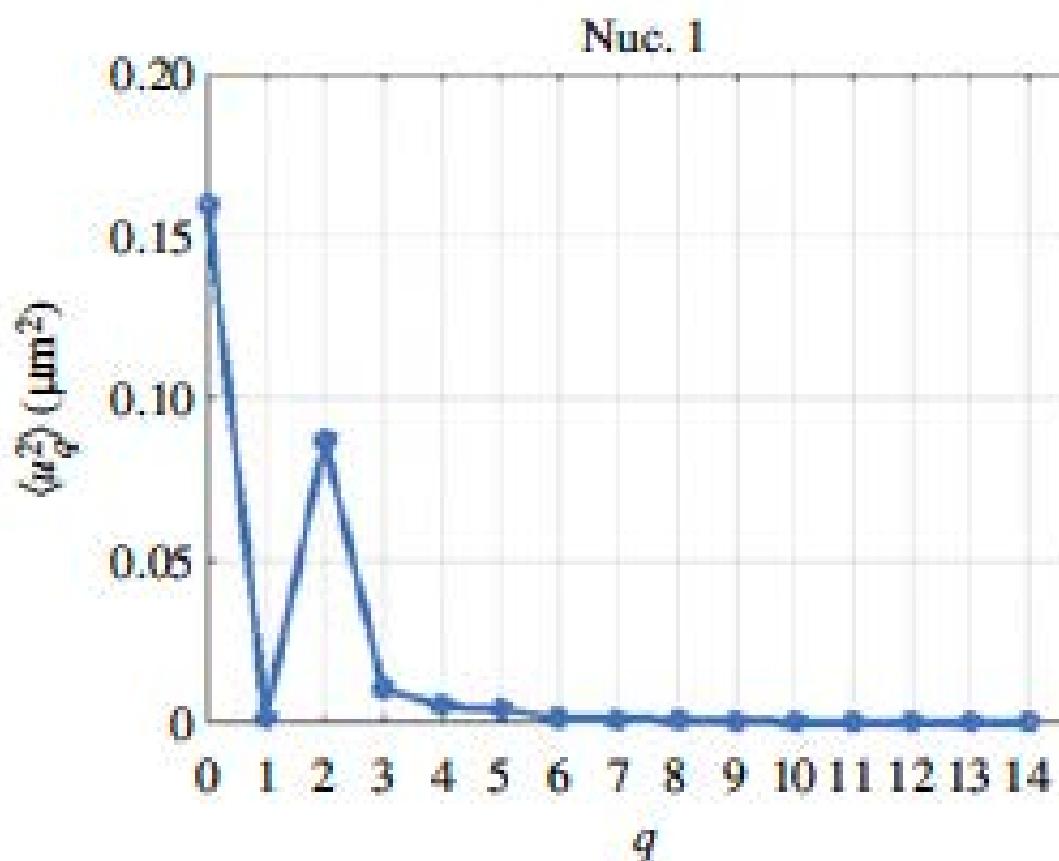
$$R_u(\delta\phi) = \frac{\langle u(\phi, t) u(\phi + \delta\phi, t) \rangle_{\phi,t}}{\langle u(\phi, t)^2 \rangle_{\phi,t}},$$

$$u(\phi, t) = r(\phi, t) - \langle r(\phi) \rangle$$

(a)



(b)



Two-time correlations

$$R_{00}(\tau) = \langle w_0(t)w_0(t+\tau) \rangle = \langle w_0(t')w_0(t'-\tau) \rangle = R_{00}(-\tau)$$

$$R_{20}(\tau) = \langle w_2(t)w_0(t+\tau) \rangle = \langle w_0(t')w_2(t'-\tau) \rangle = R_{02}(-\tau)$$

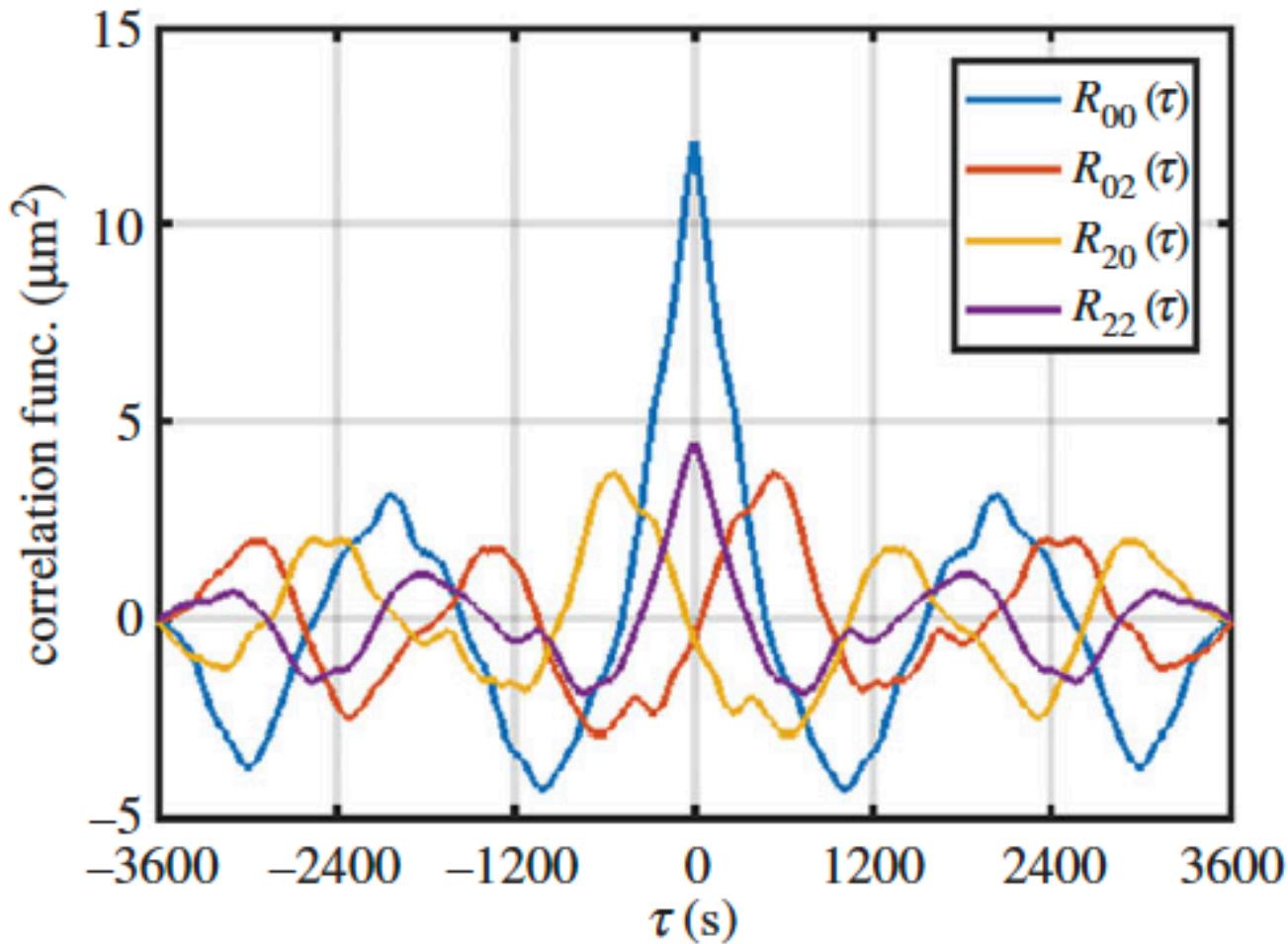
$$R_{02}(\tau) = \langle w_0(t)w_2(t+\tau) \rangle = \langle w_2(t')w_0(t'-\tau) \rangle = R_{20}(-\tau)$$

$$R_{22}(\tau) = \langle w_2(t)w_2(t+\tau) \rangle = \langle w_2(t')w_2(t'-\tau) \rangle = R_{22}(-\tau)$$

$t' = t + \tau$, time translational symmetry, product commute

If $(t, -t)$ symmetry: $R_{20}(\tau) = R_{02}(\tau)$, equilibrium

A contrario, if $R_{20}(\tau) \neq R_{02}(\tau)$: Non equilibrium



$$R_{02}(\tau) \neq R_{20}(\tau)$$

OUT OF EQUILIBRIUM

$$\frac{dw_0}{dt}(t) = A w_0(t) + B w_2(t) + \zeta_0(t)$$

$$\frac{dw_2}{dt}(t) = C w_0(t) + D w_2(t) + \zeta_2(t),$$

$$\frac{dR_{00}}{d\tau}(\tau) = A R_{00}(\tau) + B R_{20}(\tau),$$

$$\frac{dR_{02}}{d\tau}(\tau) = A R_{02}(\tau) + B R_{22}(\tau),$$

$$\frac{dR_{20}}{d\tau}(\tau) = C R_{00}(\tau) + D R_{20}(\tau)$$

$$\frac{dR_{22}}{d\tau}(\tau) = C R_{02}(\tau) + D R_{22}(\tau).$$

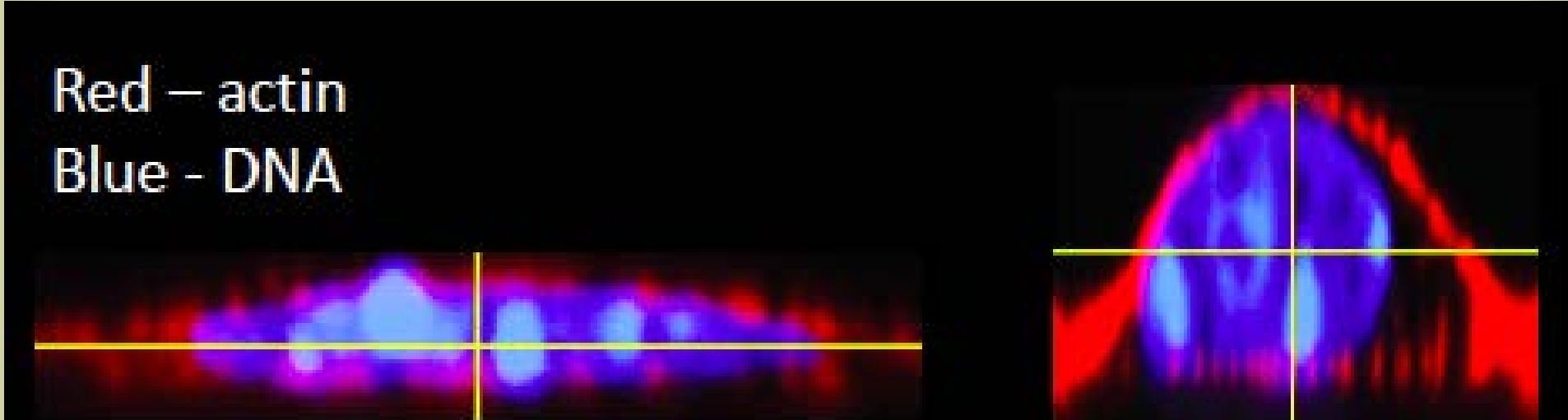
Onsager  

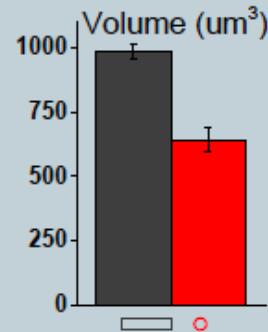
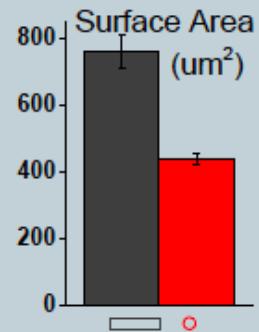
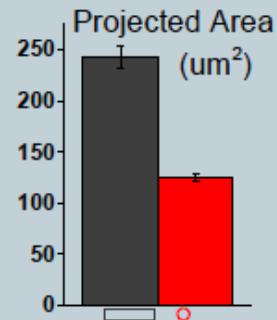
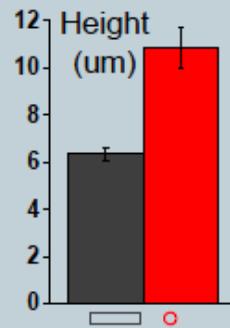
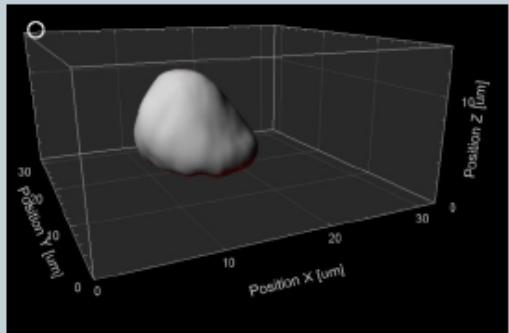
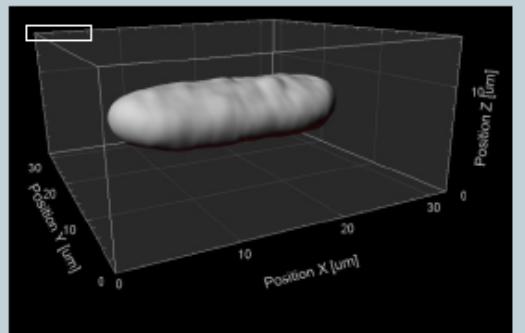
Cell Nucleus Fluctuations

differentiated cell

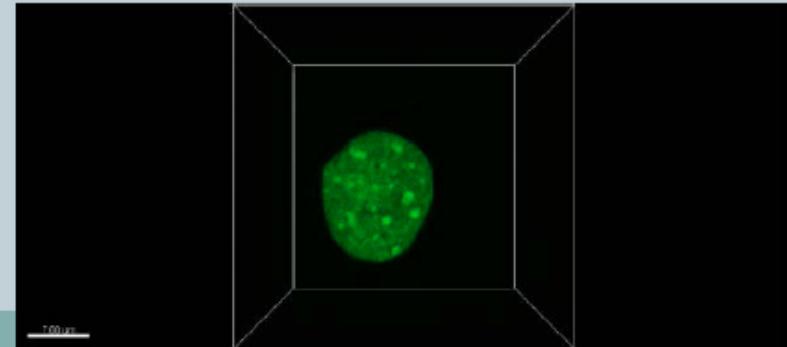
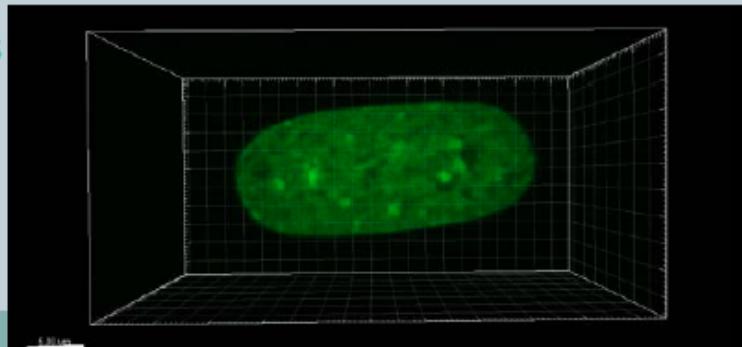
E. Makhija, D. S. Jokhun, and G. V. Shivashankar,
Proc. Natl. Acad. Sci. U.S.A. , 113 , E32 (2016).

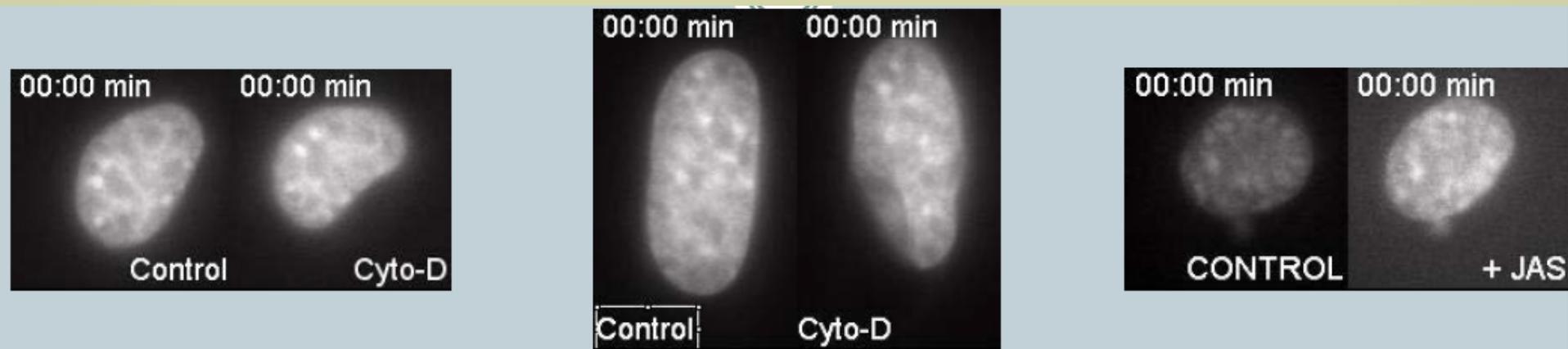
Red – actin
Blue - DNA



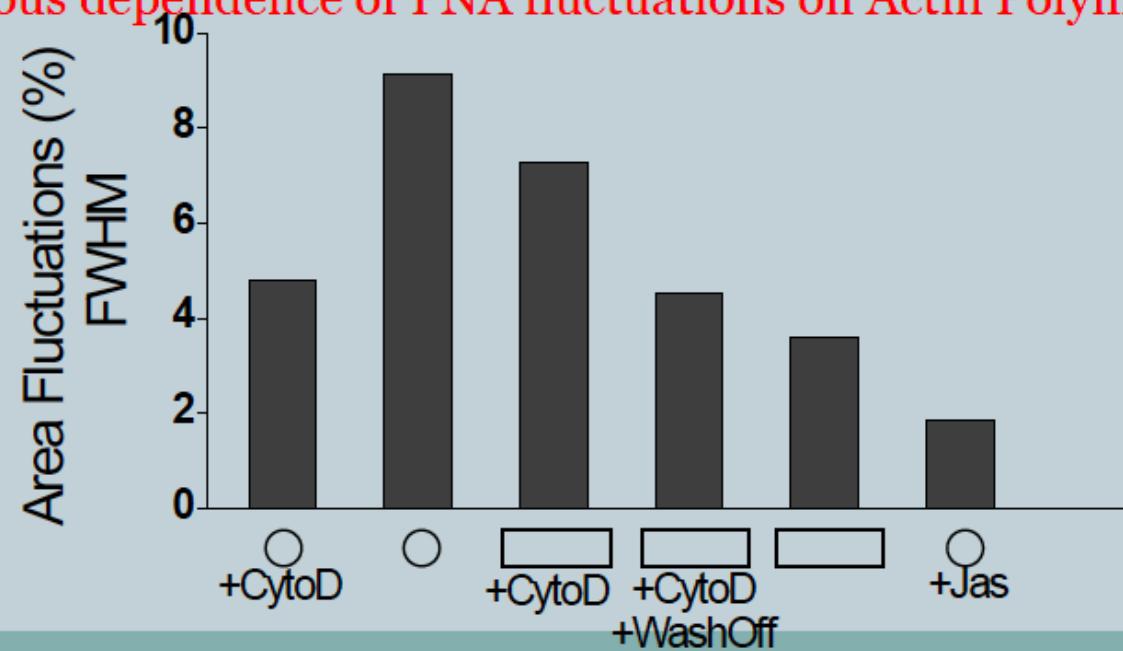


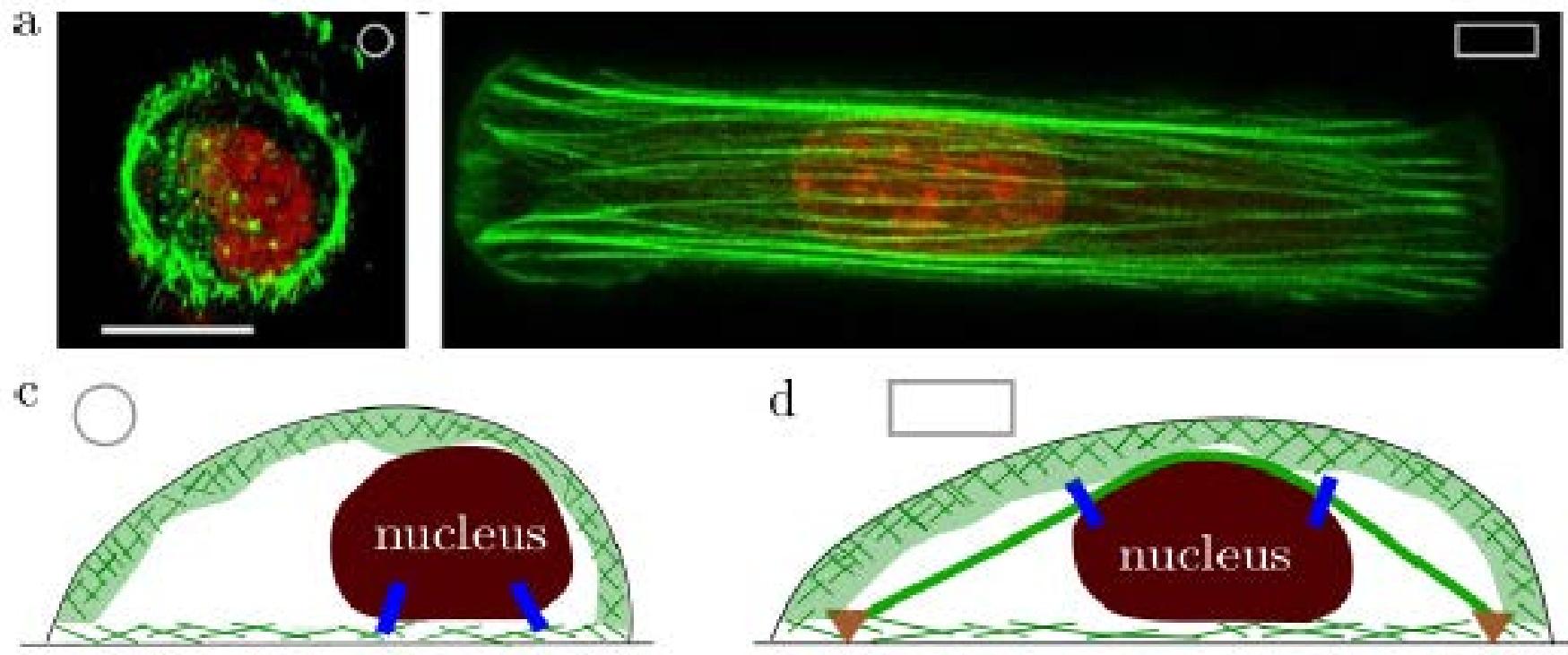
H2B





Non-monotonous dependence of PNA fluctuations on Actin Polymerization State!





J.-F. Rupprecht,¹ A. Singh Vishen,² G. V. Shivashankar,¹ M. Rao,² and J. Prost¹

$$F = -kX(t)$$

$$(\tau_v\partial_t+1)\sigma(x,t)=\eta\partial_xv(x,t)+\zeta\Delta\mu+\theta_A+\phi_T,$$

$$\langle \phi_T(x,t) \phi_T(x',t') \rangle \;=\; 2 \Lambda_T \delta_{x,x'} \delta_{t,t'}$$

$$\langle \theta_A(x,t) \theta_A(x',t') \rangle \;=\; \Lambda_A \delta_{x,x'} \exp(-|t-t'|/\tau_A)/\tau_A.$$

$$(\tau_v\partial_t+1)\sigma^{(L)}=\zeta\Delta\mu+\frac{\eta\dot{x}_t}{L+x_t}+\frac{\int_{-L}^{x_t}(\theta_A+\phi_T)}{L+x_t},$$

$$\sigma^{(L)}-\sigma^{(R)}\;=\;f(x_t)-m\dot{v}_t$$

$$\tau_v \ddot{v}_t + \dot{v}_t + \frac{\lambda(x_t)}{m} v_t = \frac{f + \tau_v \dot{f} + \mu(x_t) \left[\Theta_A + \Phi_T \right]}{m},$$

$$(\langle\Theta_A(t)\Theta_A(s)\rangle ~=~ \Lambda_A\exp(-|t~-~s|/\tau_A)/\tau_A)$$

$$\langle \Theta_A(t)\Theta_A(s)\rangle ~\equiv~ \Lambda_A\exp(-|t~-~s|/\tau_A)/\tau_A)$$

$$\mu^2(x)=\frac{2L}{L^2-x^2}=\frac{\lambda(x)}{\eta}.$$

$$Multiplicative\;noise$$

$$\tau_v ~\ll~ \tau_A ~\ll~ t_u, \qquad \qquad t_u ~=~ \eta^2 L / \max(\Lambda_A,\Lambda_T)$$

$$\dot{x}_t = \frac{f}{\lambda} - \frac{\Lambda_T \mu \mu'}{2 \lambda^2} + \frac{\mu \left[\Theta_A + \Theta_T \right]}{\lambda},$$

$$\dot{x}_t = \frac{f}{\lambda} - \frac{\Lambda_T \mu \mu'}{2 \lambda^2} + \frac{\mu \left[\Theta_A + \Theta_T \right]}{\lambda},$$

$$\dot{x}_t = \frac{f}{\lambda} - \frac{\Lambda_T \mu \mu'}{2 \lambda^2} + \frac{\mu \left[\Theta_A + \Theta_T \right]}{\lambda},$$

$$\dot{x}_t = \frac{f}{\lambda} - \frac{\Lambda_T \mu \mu'}{2 \lambda^2} + \frac{\mu \left[\Theta_A + \Theta_T \right]}{\lambda},$$

$$\partial_t P = \partial_x\left\{\left[-\frac{f}{\lambda} + \frac{\Lambda_T\mu\mu'}{2\lambda^2}\right]P + \frac{\Lambda\mu}{2\lambda}\partial_x\frac{\mu}{\lambda}P\right\},$$

$$= \Lambda = (\Lambda_T^2 + \Lambda_A^2)^{1/2}.$$

$$\partial_t P = \partial_x\left\{-\frac{f}{\lambda}P + \Lambda_A\frac{\mu}{2\lambda}\partial_x\frac{\mu}{\lambda}P\right\}$$

$$P(x)=\frac{\sqrt{\pi}\Gamma\left(k+\frac{1}{2}\right)}{L\Gamma(k+1)}(1-(x/L)^2)^{-1/2+k},$$

$$k = (KL^2\eta)/(2\Lambda_A)$$

$$\partial_t P = \partial_x\left\{\left[-\frac{f}{\lambda} + \frac{\Lambda_T\mu\mu'}{2\lambda^2}\right]P + \frac{\Lambda\mu}{2\lambda}\partial_x\frac{\mu}{\lambda}P\right\},$$

$$= \Lambda = (\Lambda_T^2 + \Lambda_A^2)^{1/2}.$$

Maximal fluctuations when noise-induced destabilization and focusing force are balanced

$$P(x) = \frac{\sqrt{\pi}\Gamma\left(k + \frac{1}{2}\right)}{\Gamma(k+1)}(1-x^2)^{-1/2+k}, \rightarrow \text{Var}[X] \propto 1/(k+1/2)^{3/2}$$

no optimal restoring force/activity?

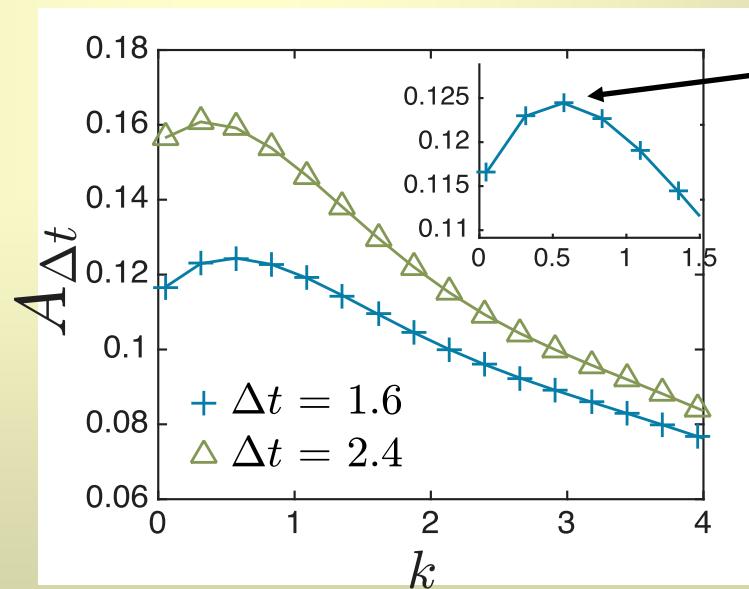
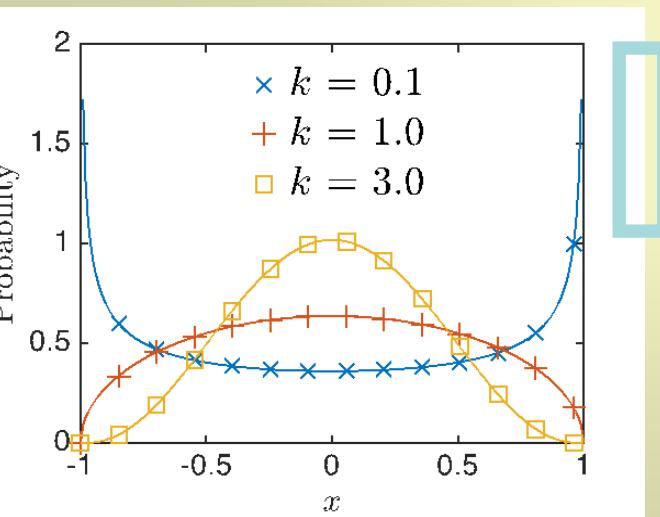
> what is measured in experiments is

$$A_{\Delta t} = \langle (X - \bar{X}_{\Delta t})^2 \rangle_s$$

$\langle . \rangle_s$: average over large number of cells (random realisation)
 $\bar{X}_{\Delta t}$ is an average position over an observation window Δt

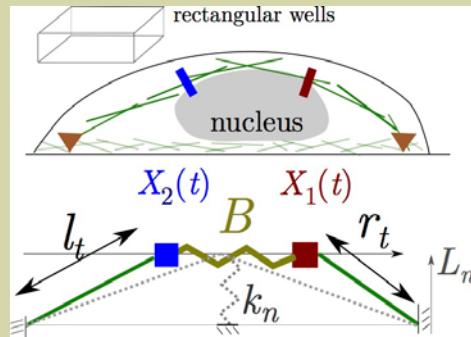
Mean time to explore the whole gel: $t_{\text{MFPT}} \approx 10 \times \eta/\gamma$

short-times ($\Delta t < t_{\text{MFPT}}$): exploration limited to maximal $P(x)$ regions



$k_{\text{opt}} \rightarrow 1/2$

long-times ($\Delta t > t_{\text{MFPT}}$): $A_{\Delta t} \rightarrow \text{Var}[X] \propto 1/K \rightarrow k_{\text{opt}} \rightarrow 0$

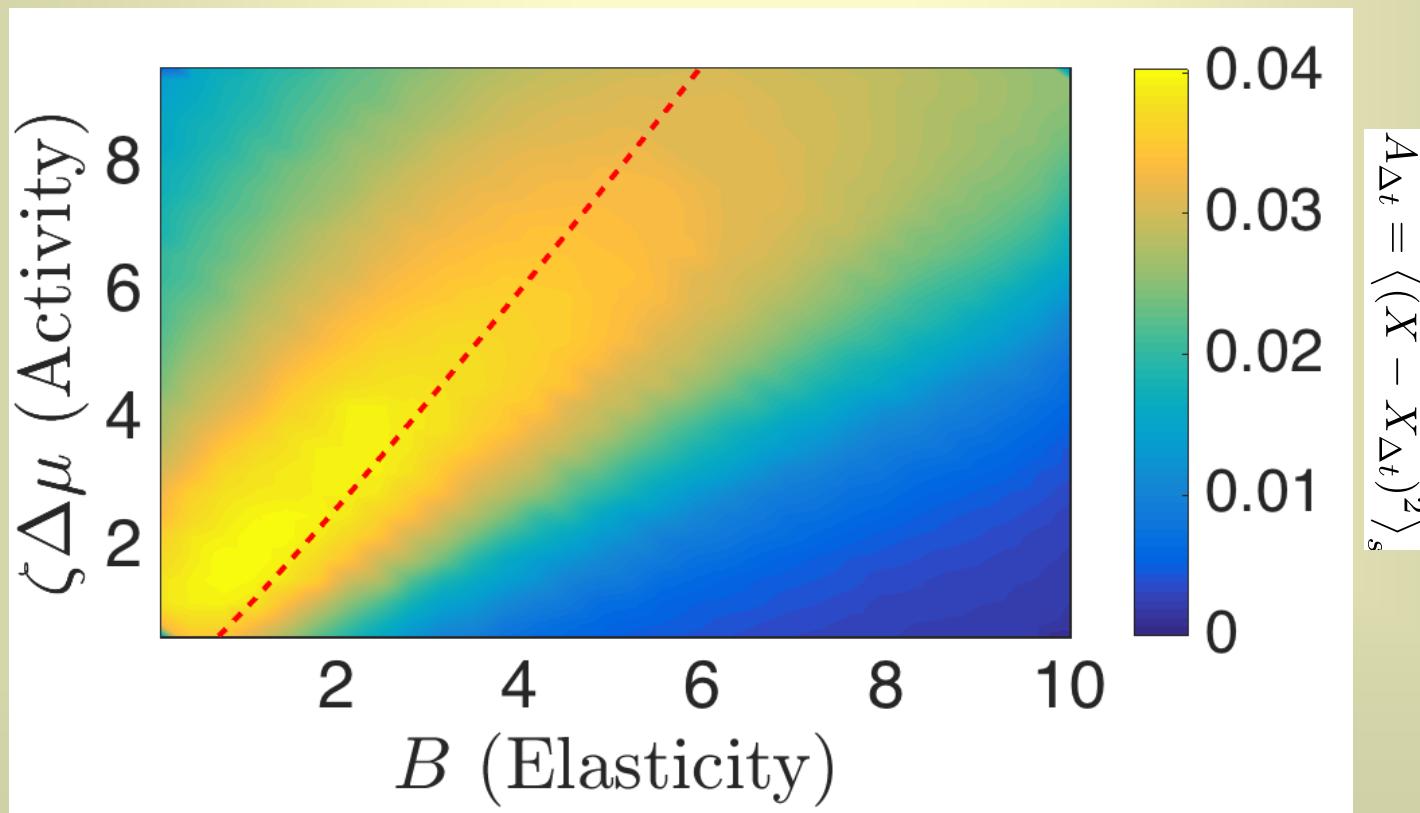


Limit case:

$$L_n = 0$$

$$\zeta \Delta \mu_{opt} \square 2B - \sqrt{B}$$

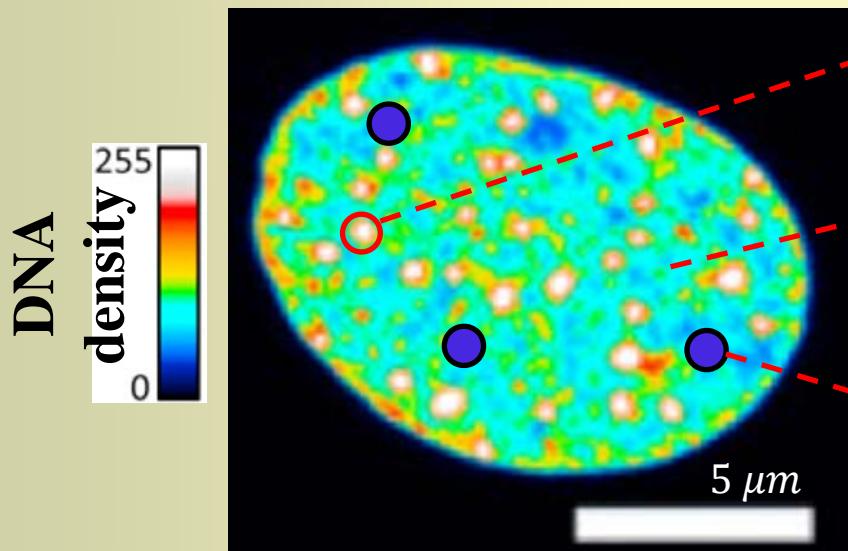
Optimum Activity
exist again!



How enzymatic activity is involved in chromatin organization

Rakesh Das , Takahiro Sakaue, GV Shivashankar, Jacques Prost, Tetsuya Hiraiwa

➤ Phase separated organization of chromatin



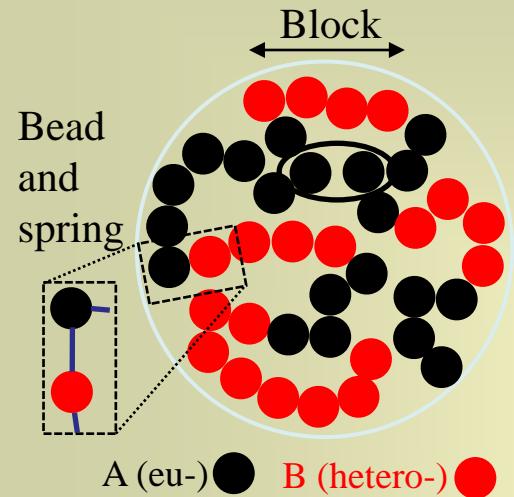
Heterochromatin –
transcriptionally repressive

Euchromatin –
transcriptionally active

[Transcription – copying DNA into RNA]

Subnuclear condensates (SNCs)
—
droplets of biochemical agents
(schematic)

Copolymer model with Transient-Linking Activity (TLA)

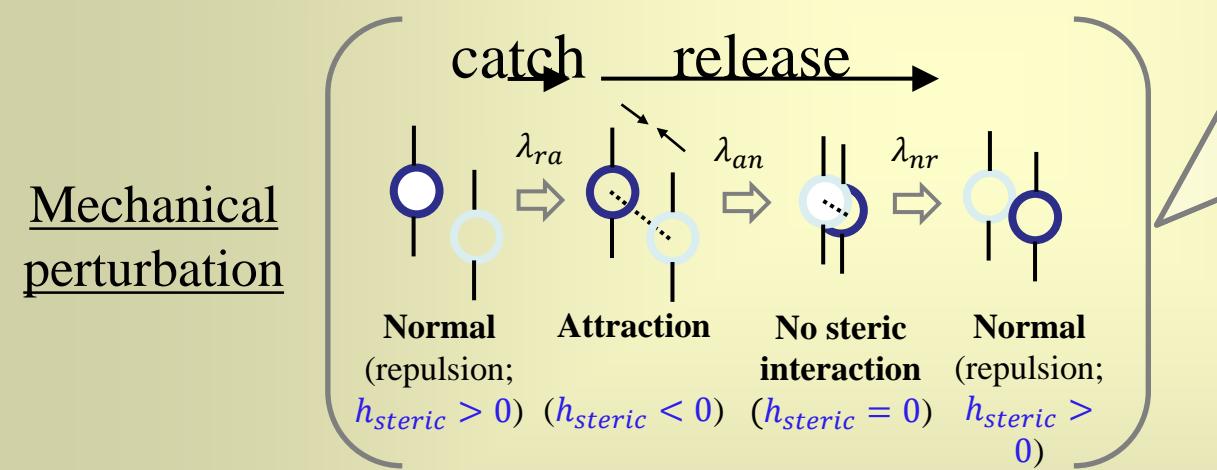


$$\frac{\partial \mathbf{x}_b}{\partial t} = -\Gamma \frac{\partial H(\mathbf{x}_b)}{\partial \mathbf{x}_b} + \sqrt{2\Gamma} \boldsymbol{\zeta}_b$$

$$H = \sum_{n. n.} h_{chain} + \sum_{\text{short range}} h_{steric} - \sum_{\text{short range}} h_{HC \text{ affinity}} + H_{confine.}$$

(short range)

$\boldsymbol{\zeta}_b \equiv$ Univariate white Gaussian noise with zero mean



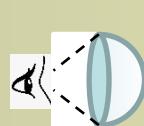
Reminiscent to Topoisomerase-II enzyme's action – resolves topological constraints of chromatin



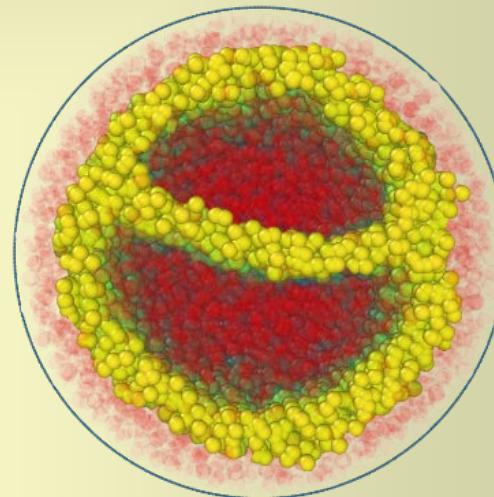
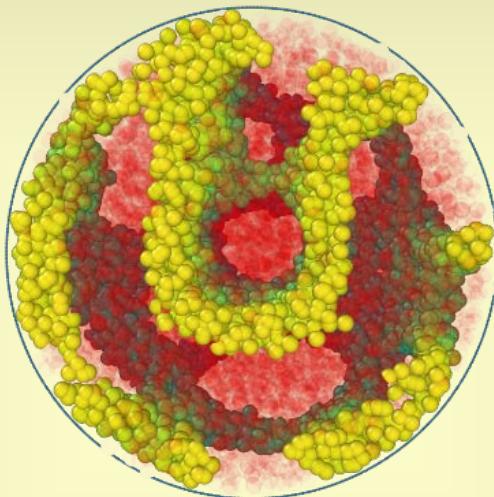
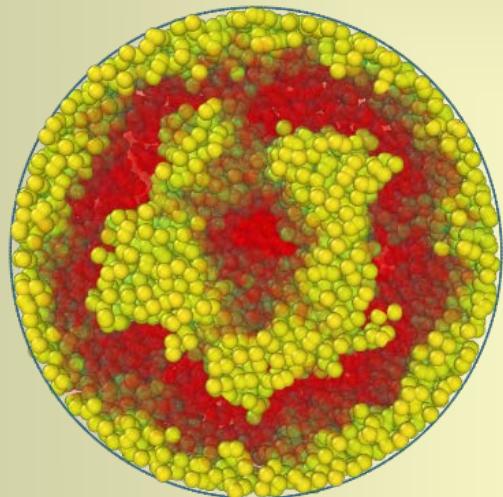
J. Roca, Nucleic Acids Res. 37, 721 (2009)

- Transient-linking activity (TLA) = $\frac{\text{rate of catching}}{\text{rate of releasing}} = \lambda_{ra} \left(\frac{1}{\lambda_{an}} + \frac{1}{\lambda_{nr}} \right)$
- Only AA (i.e., eu-eu) pairs are subjected to TLA [A. S. Sperling et al., PNAS 108, 12693 (2011)]

Transient-linking activity (TLA) alters chromatin organization



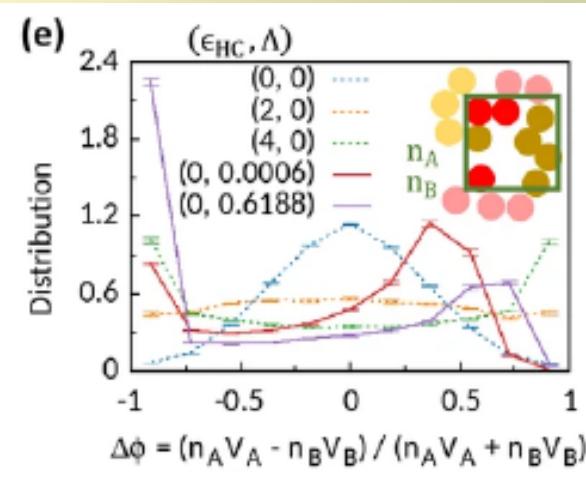
↓ + TLA → Time



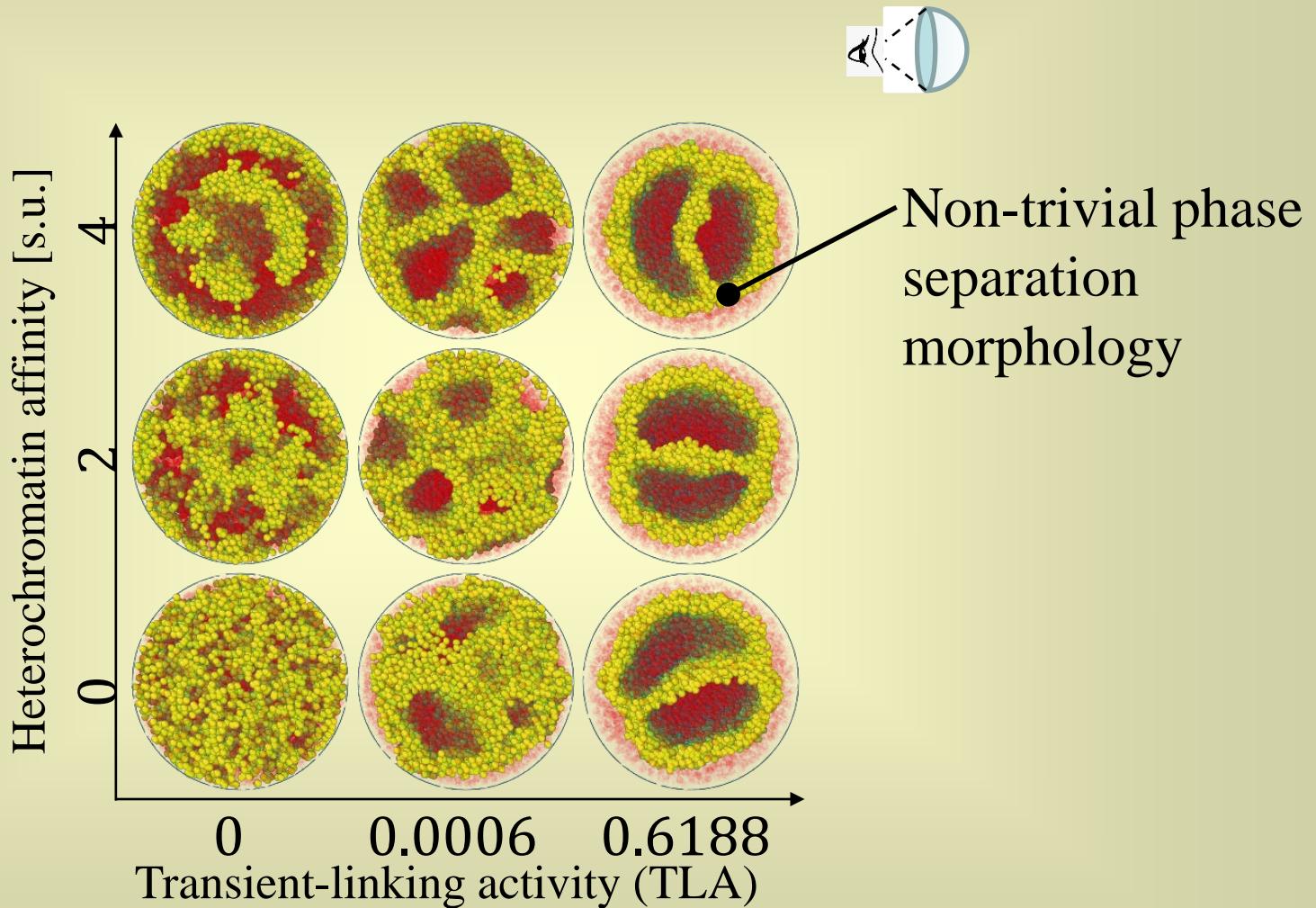
Heterochromatin
affinity

Euchromatin: Pole Z-coord. of Equator

Heterochromatin: semi-transparent red



List of typical snapshots in transient-linking activity (TLA) vs. heterochromatin affinity phase space



Happy Birthday!

