

# modeling the hysteretic behavior of chromatin fibers under magnetic tweezers:

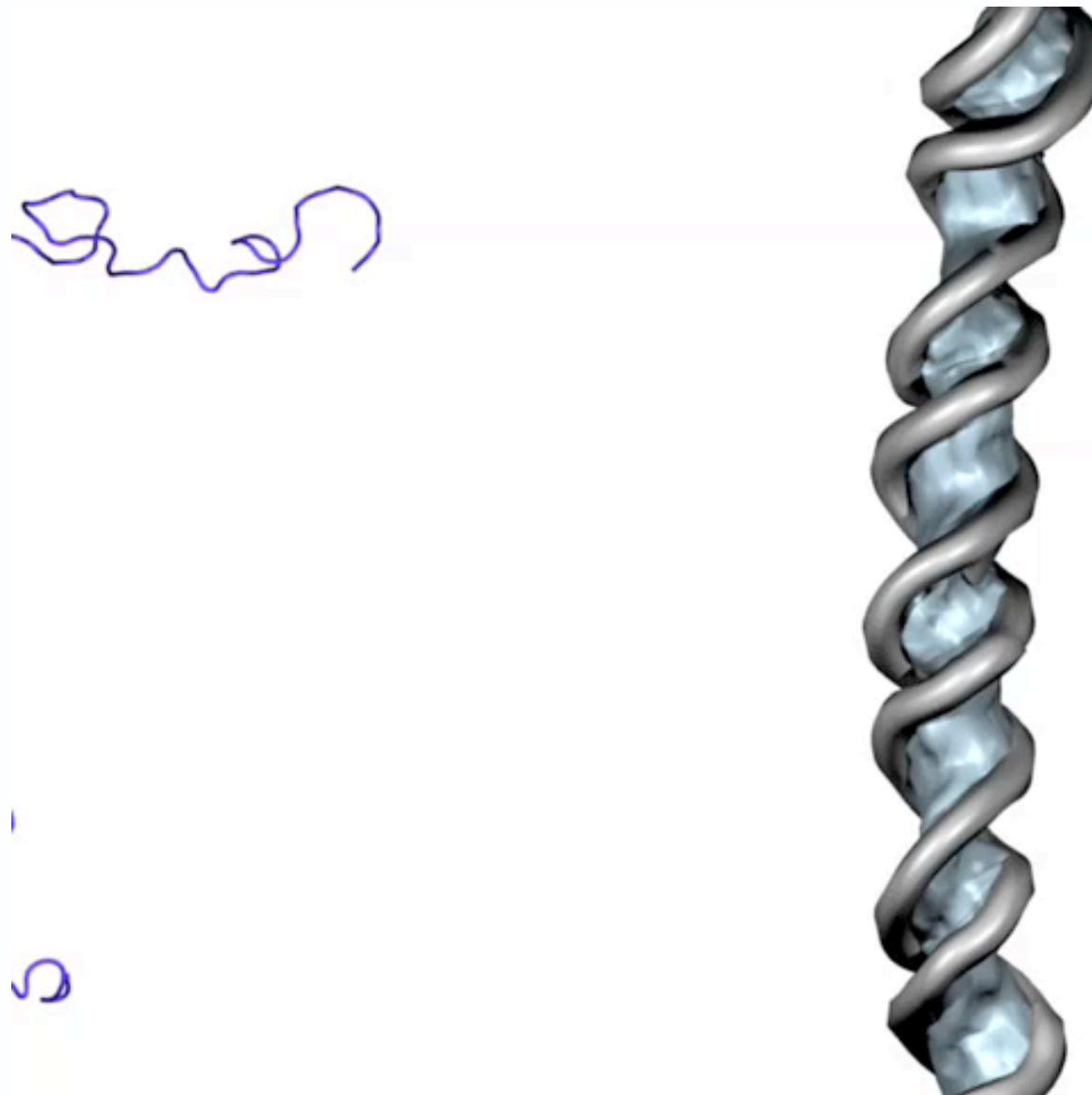
*why  
are nucleosomes  
left handed?*

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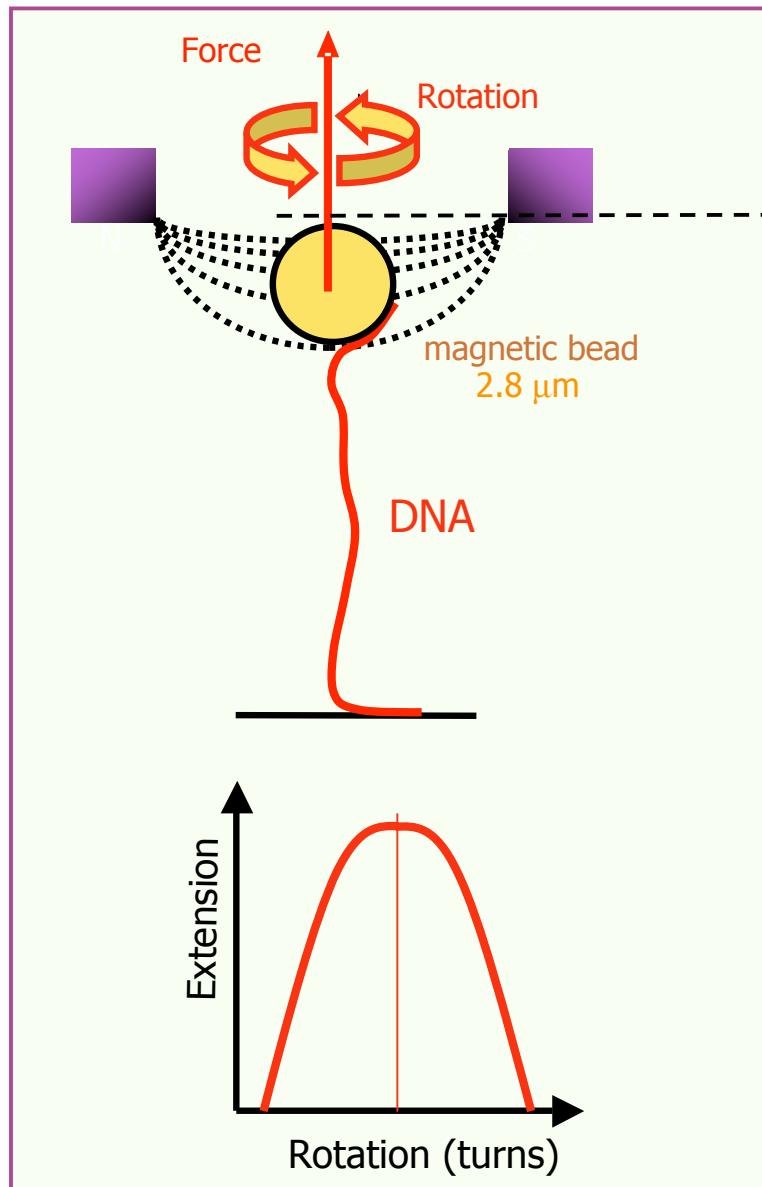
*experimental group:  
Jean-Louis Viovy, Institut Curie, Paris*

# nucleosome structure

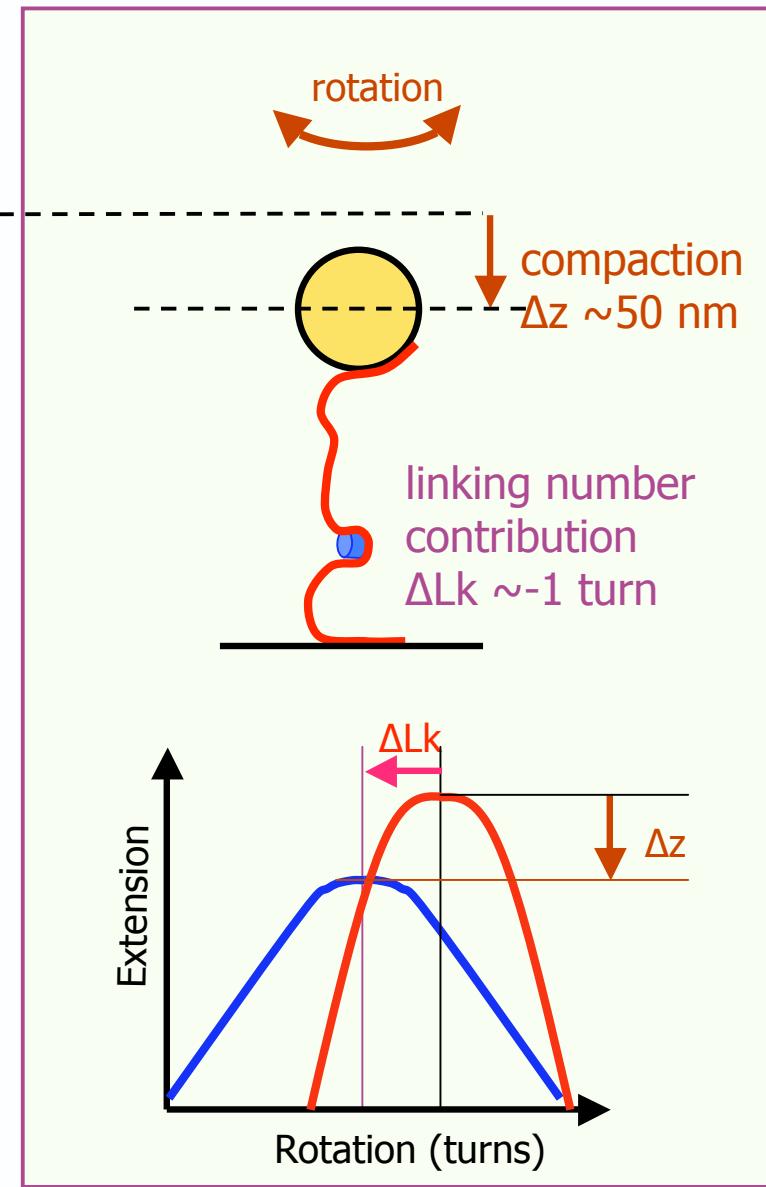


Animation by Hua Wong

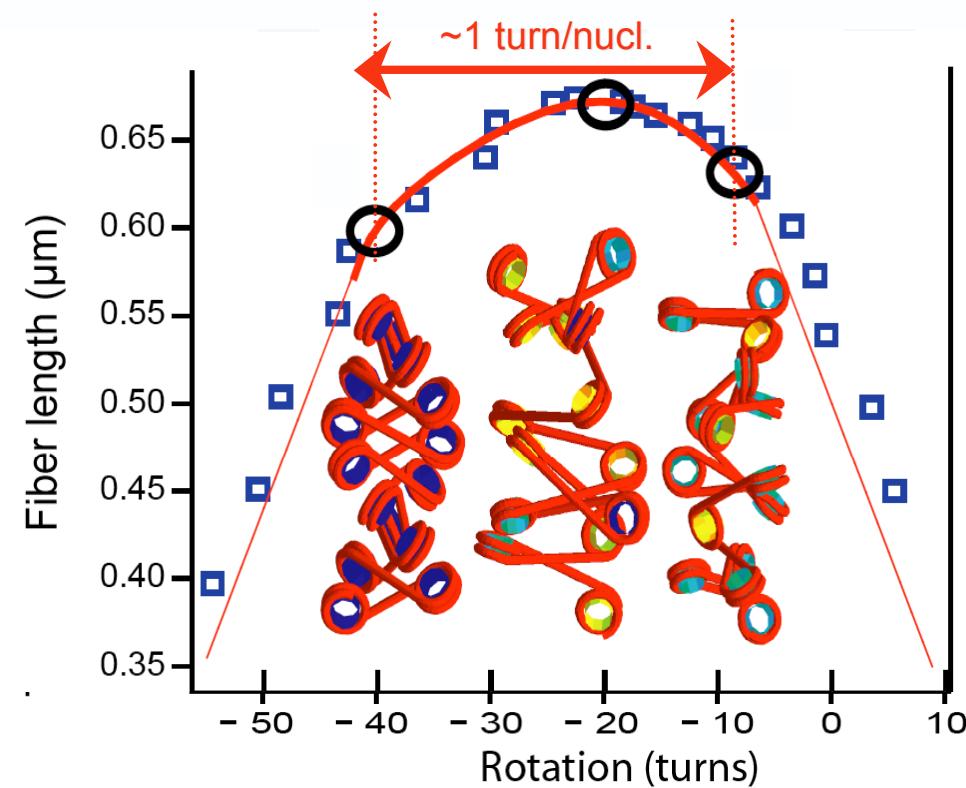
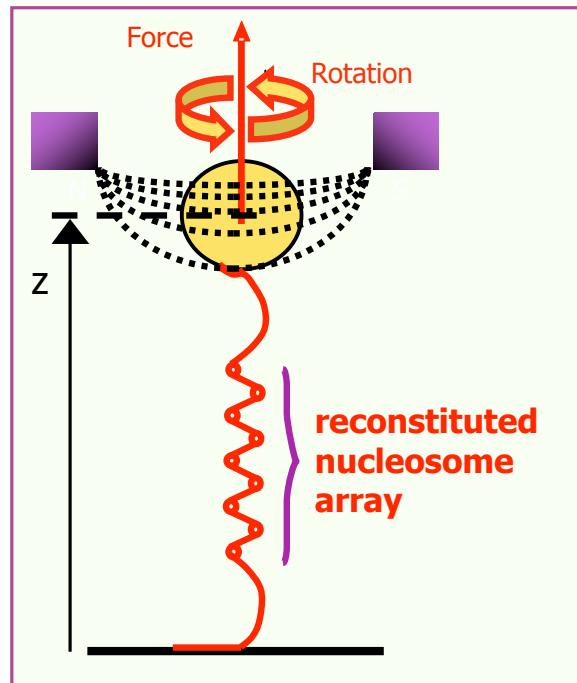
# probing structural features with magnetic tweezers: nucleosome



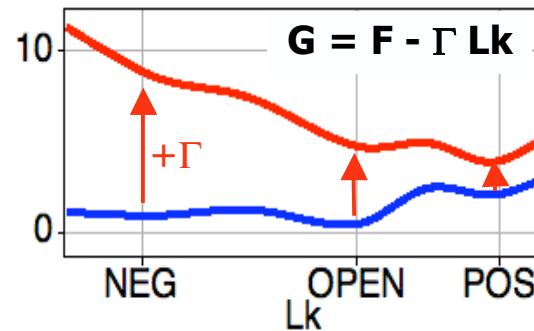
+ 1 nucleosome



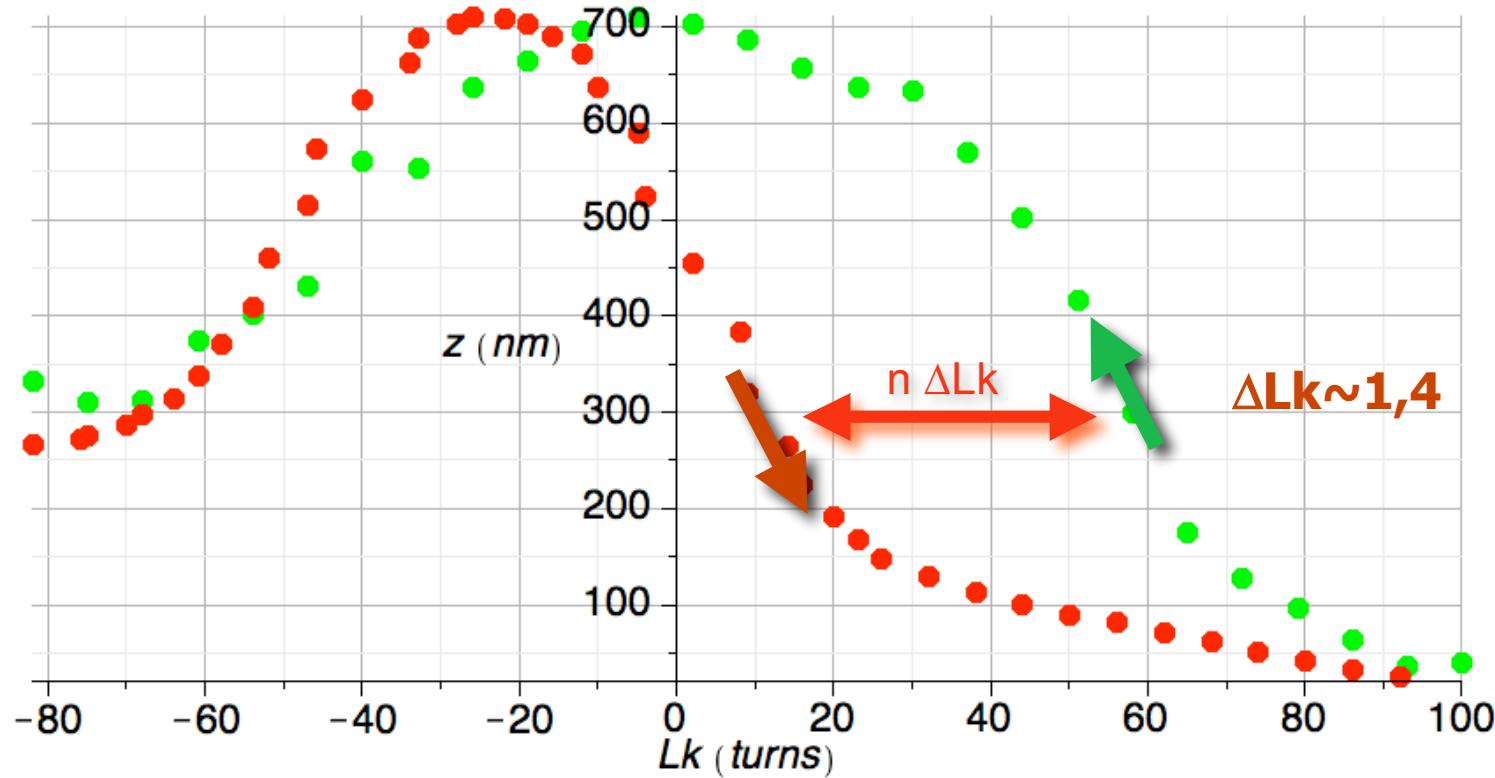
# probing structural features with magnetic tweezers: **FIBER**



**Model:**  
torque-dependent equilibrium between the 3 states of nucleosome



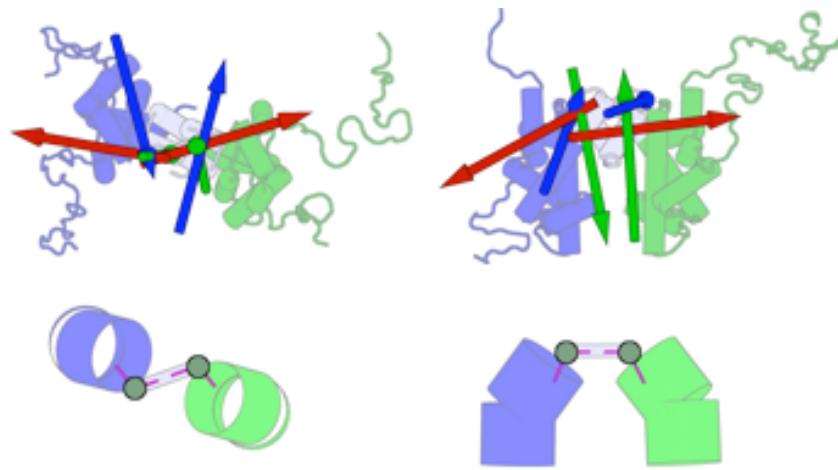
# hysteretic response under extensive torque



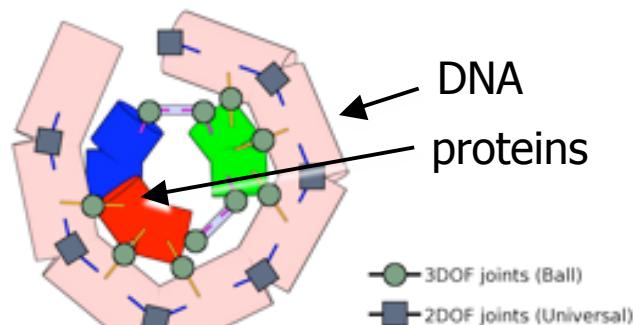
- not explained by nucleosome-nucleosome interactions
- compatible with a **nucleosome internal rearrangement: transition to a metastable altered state with a larger (positive)  $\Delta Lk$**

# Rigid Body Dynamics with Skeletal Animation

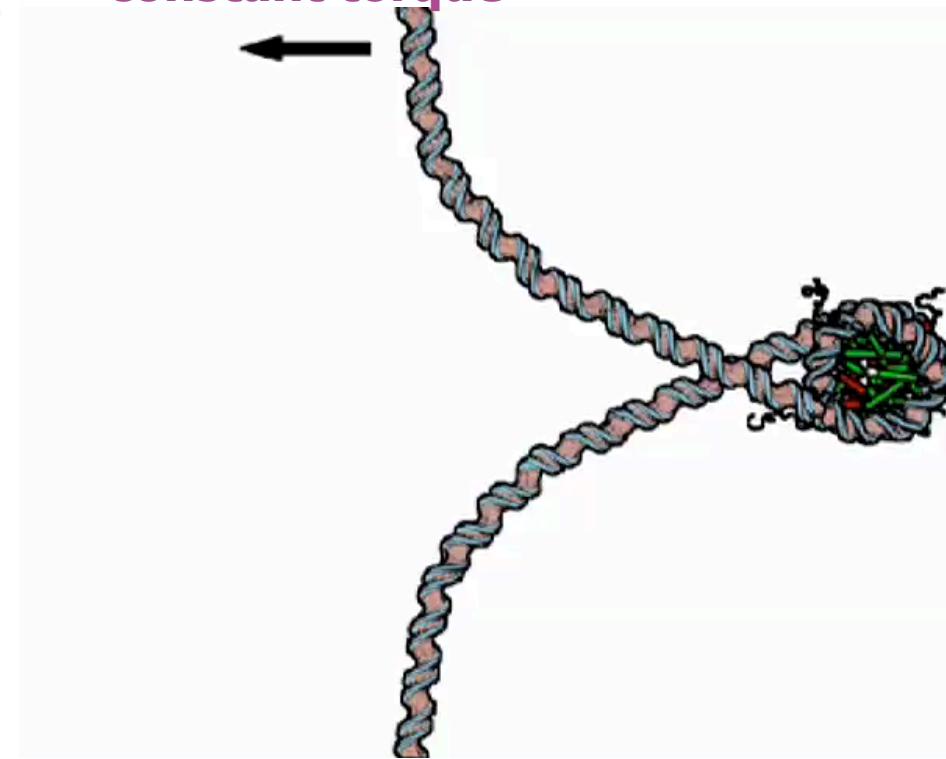
## 1. normal modes analysis



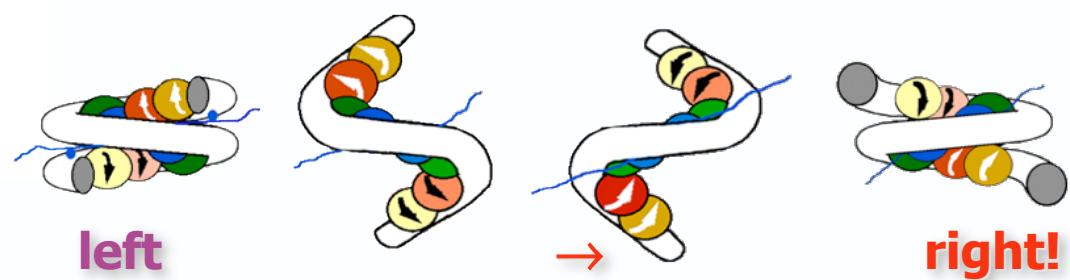
## 2. nucleosome structuration



## 3. Brownian dynamics under constant torque



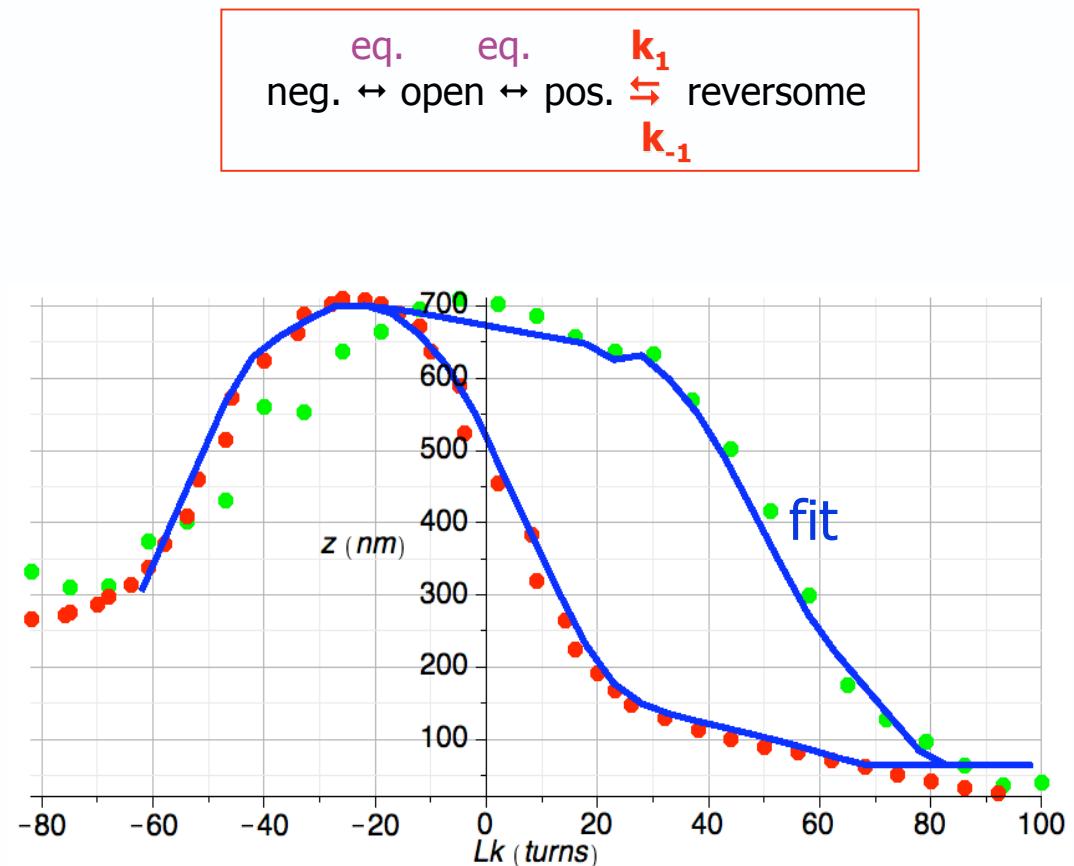
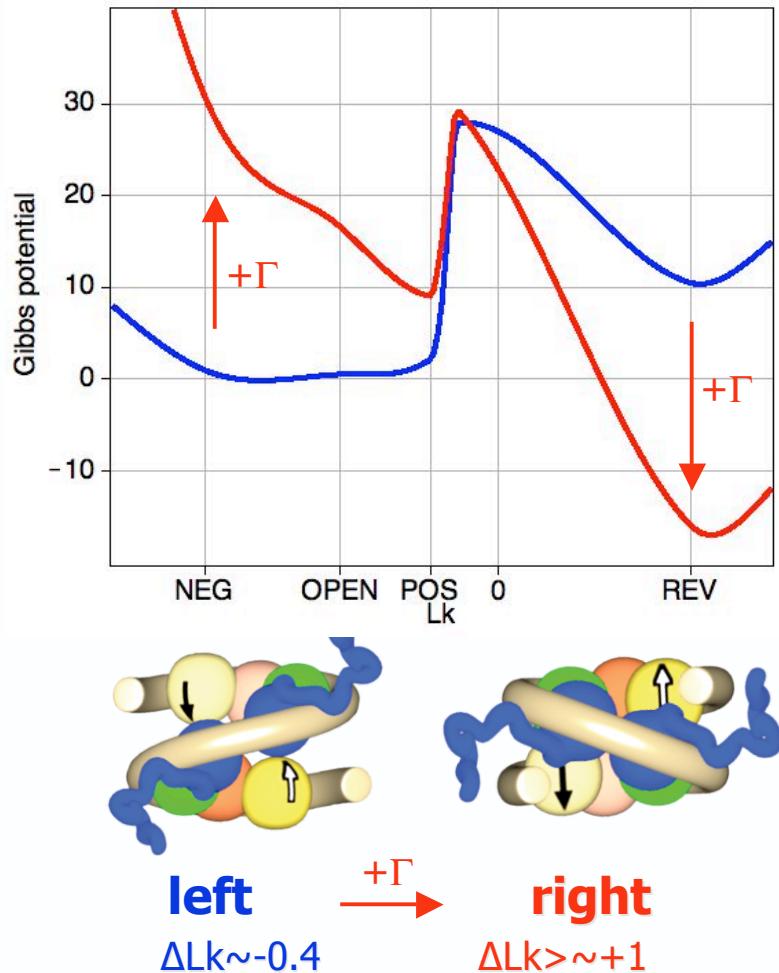
**result: chiral transition:**



Animation by Hua Wong

# **Reversome : a right handed metastable state... stabilized by the applied torque**

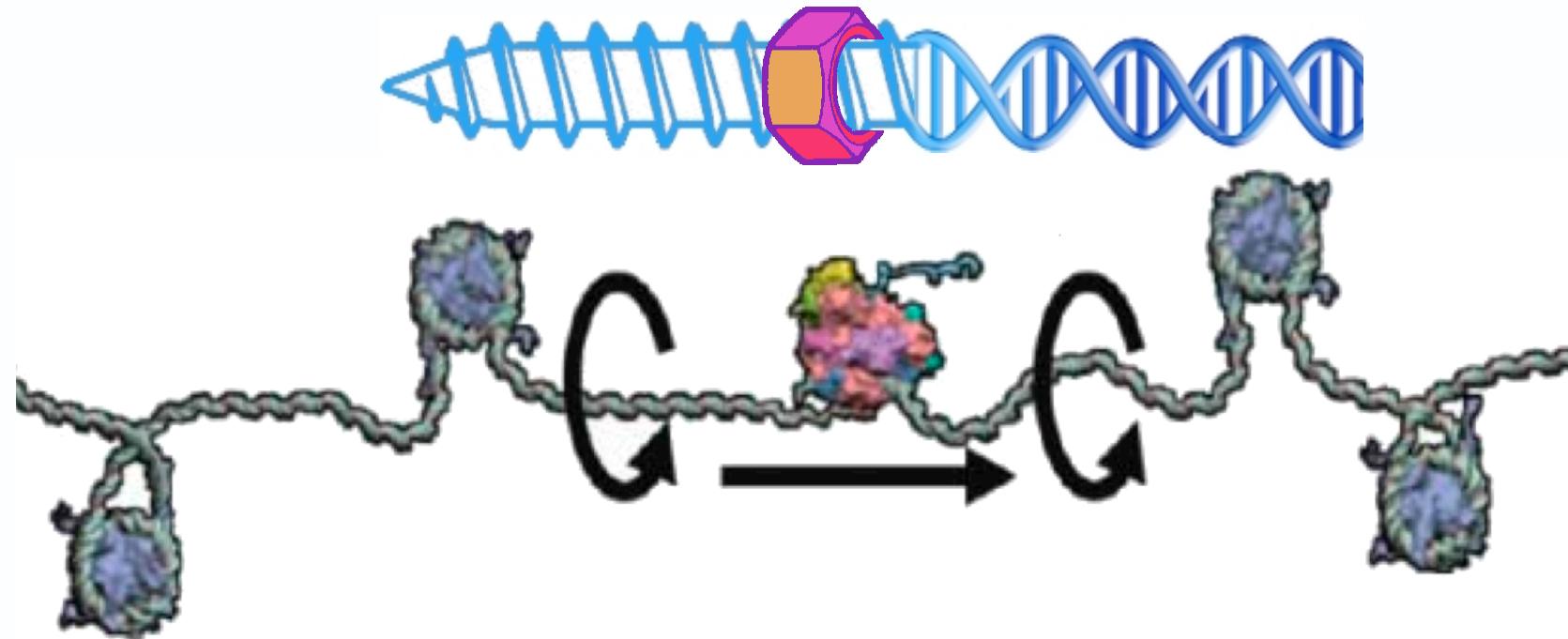
data fit: statistical equilibrium between the 3 states  
+ kinetic equations for the passage to reversome:



So, why *standard* nucleosome is left handed ?

*In vivo* prediction:

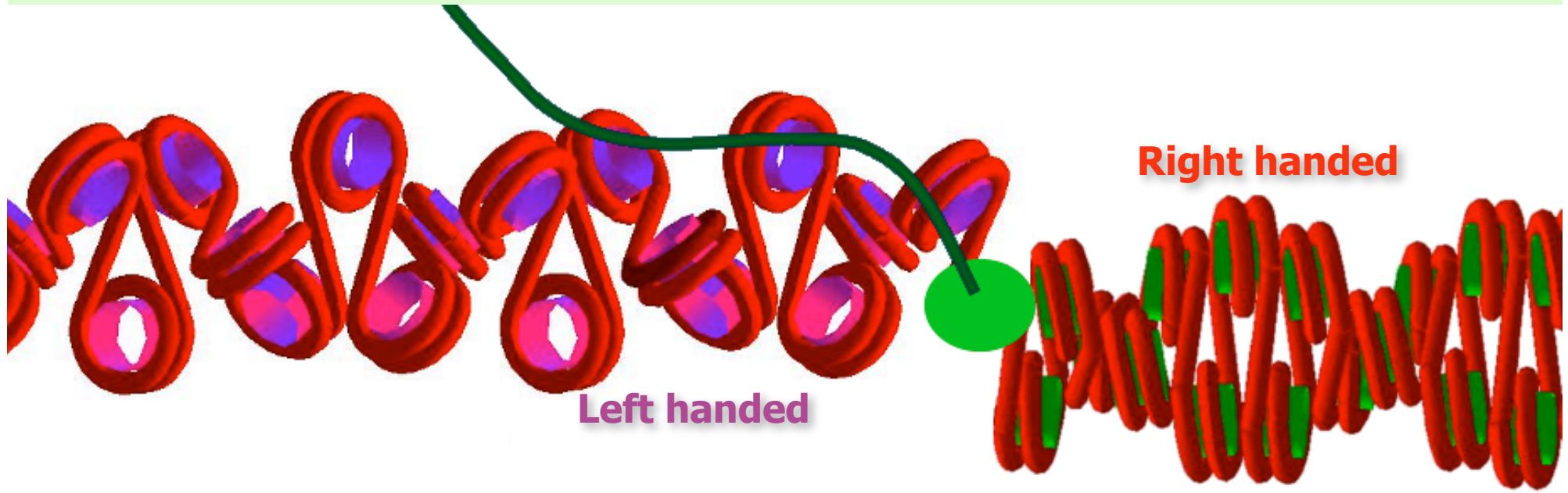
- RNA-polymerase fixed in the transcription complex
- right handed DNA screws through the polymerase



⇒ RNA polymerase exerts a pulling force and a positive torque on DNA

# So, why *standard* nucleosome is left handed ?

***Conclusion: a possible functional role for nucleosome:***



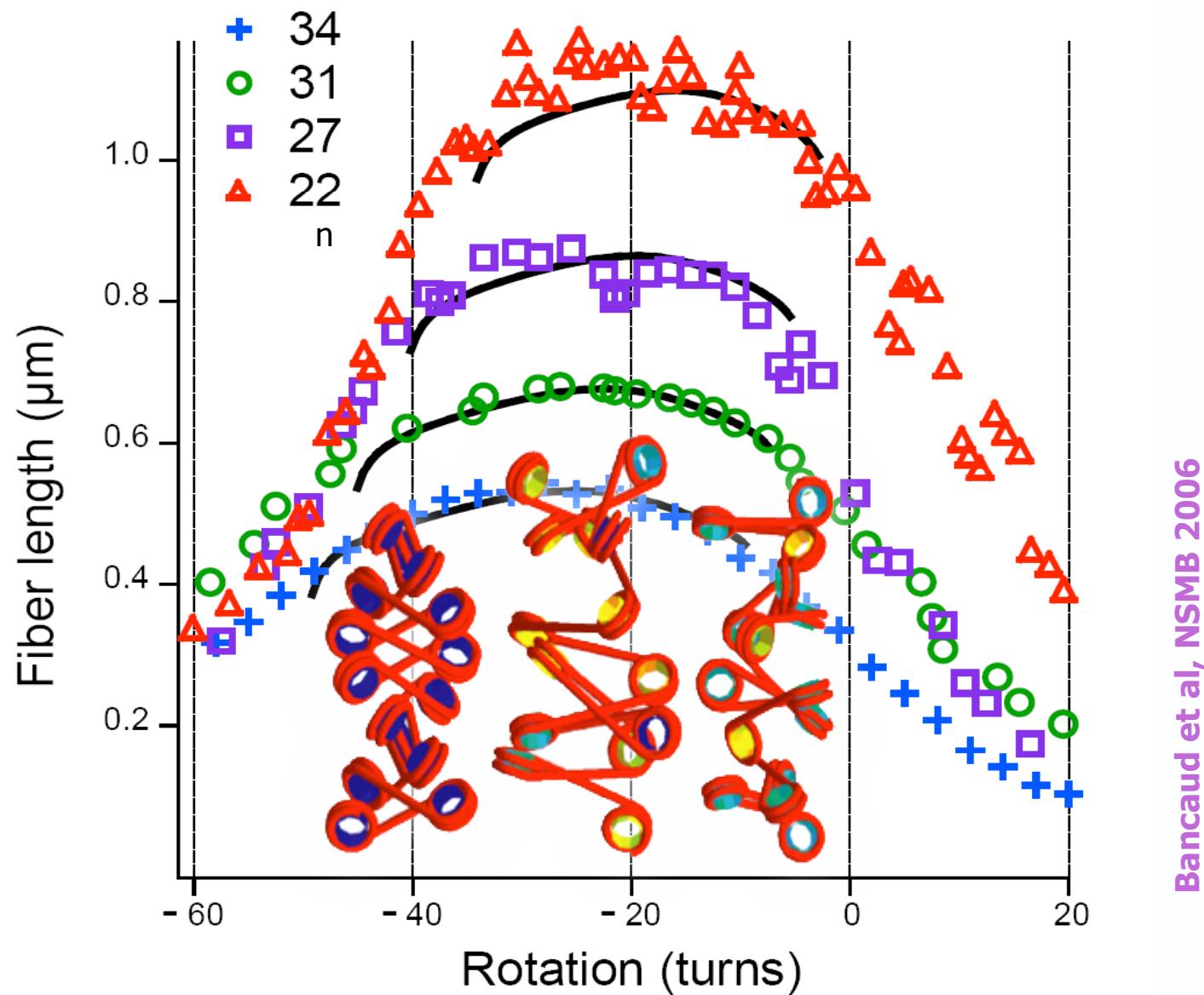
- nucleosome → reversome transition absorbs turns and thus releases supercoiling constraint
- reversome is a destabilized state favoring the passage of the RNA-polymerase



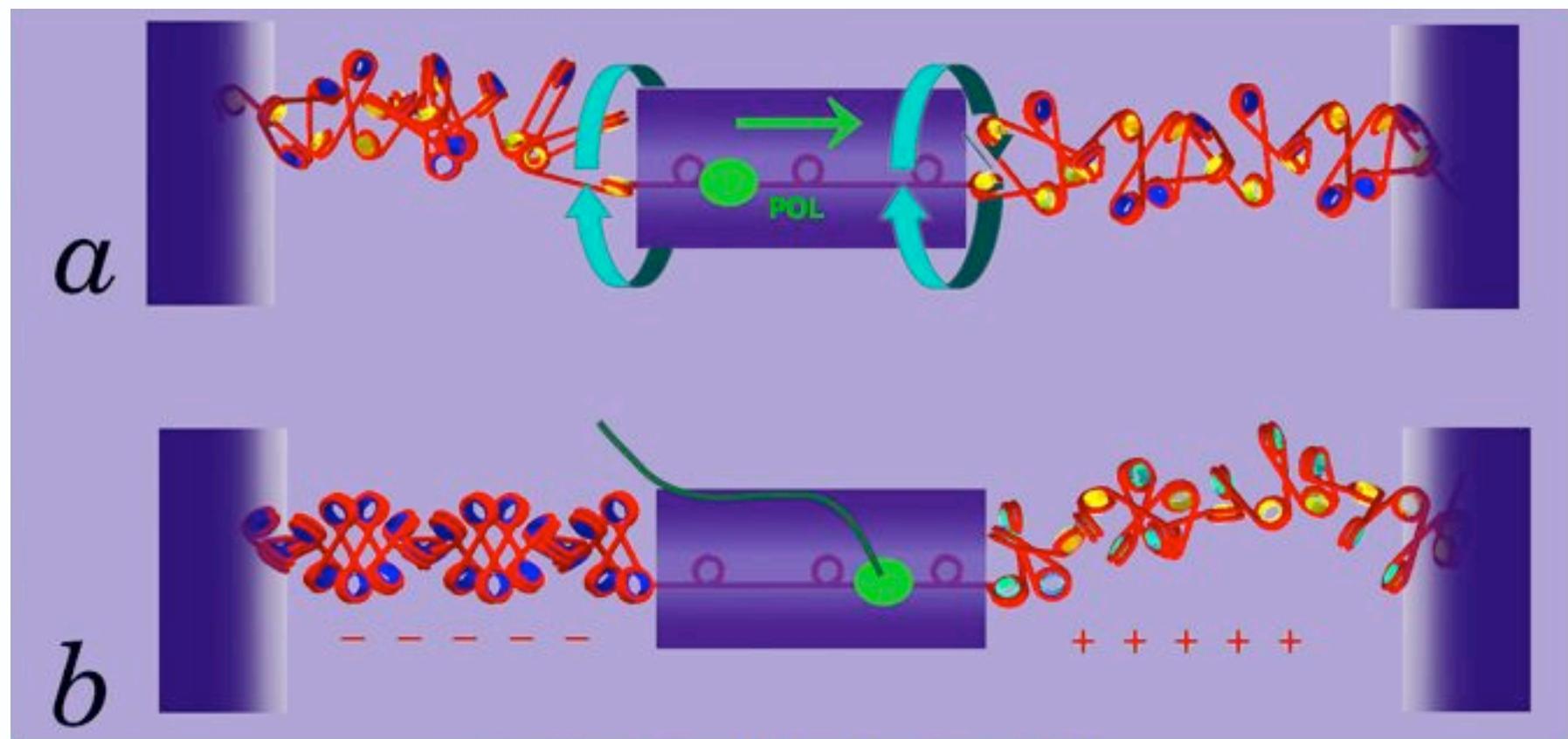
# M3V Group

- **Maria Barbi** (UPMC)
- **Annick Lesne** (CNRS)
- **Jean-Marc Victor** (CNRS)
- **Julien Mozziconacci** (doc 2001-2004,  
postdoc ANR 2006–2008)
- **Hua Wong** (doc 2005-2008)
- **Fabien Paillusson** (doc 2007-2010)
- **Christophe Becavin** (doc 2007-2010)
- **Christophe Lavelle** (postdoc 2003-2004)

# Experimental data fit for different numbers of nucleosomes

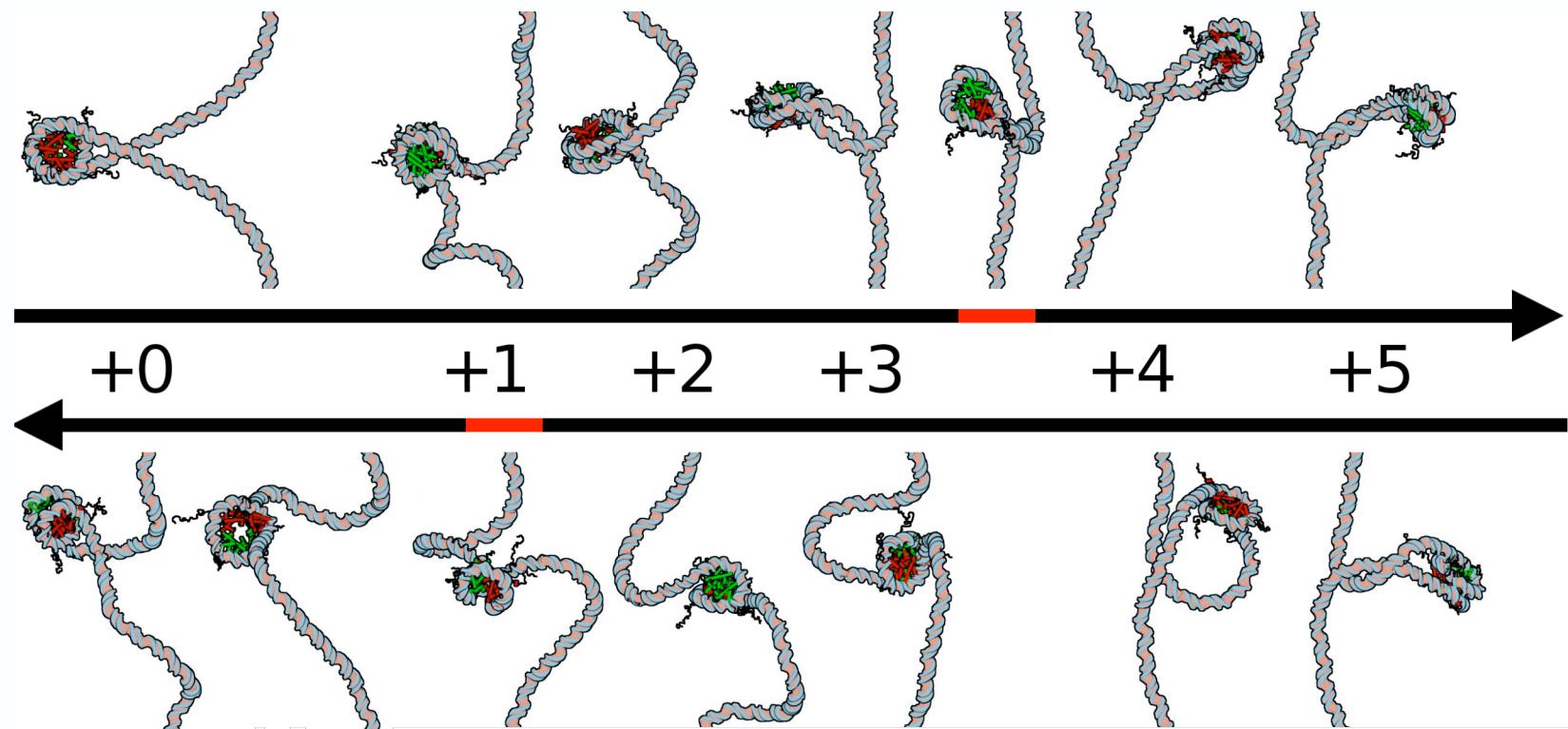


# The three states acts as a topological buffer



(Bancaud et al, NSMB 2006)

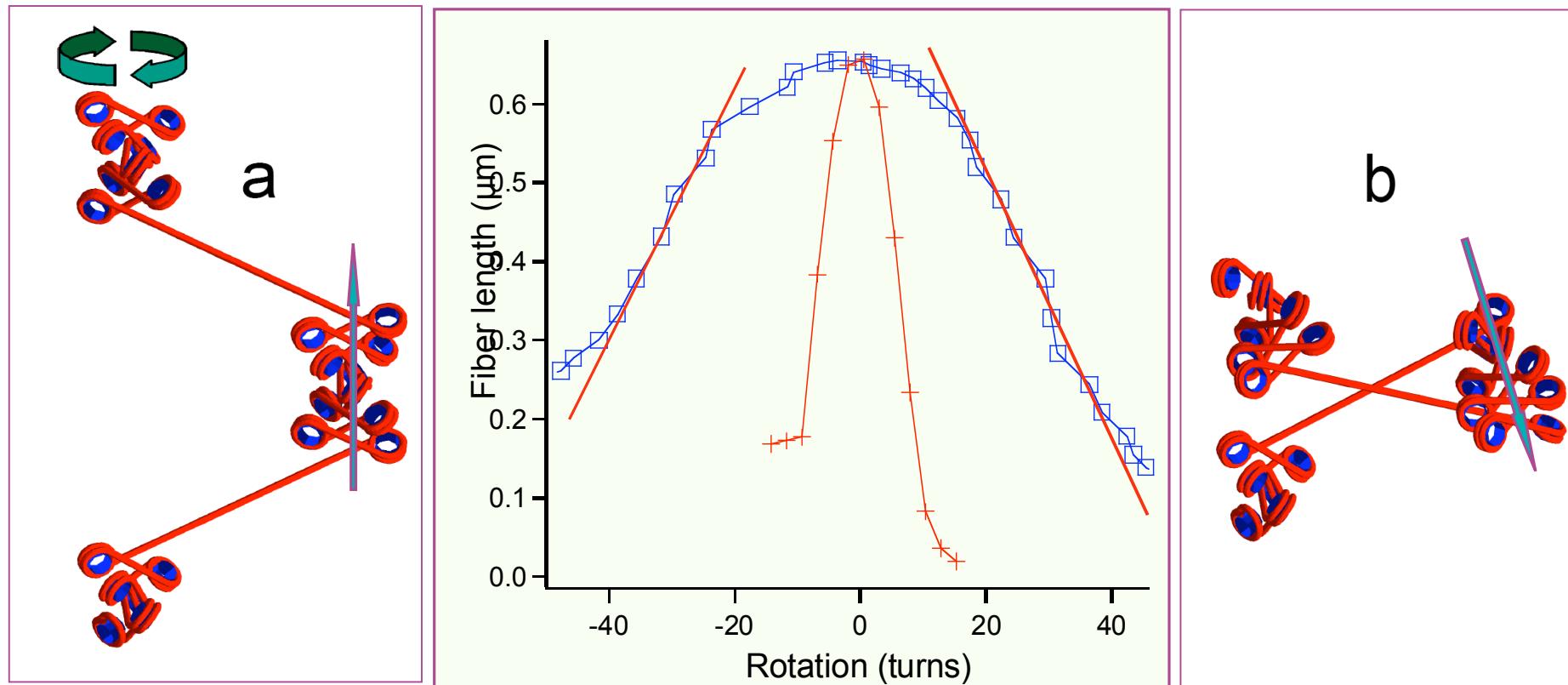
# Looking at the transition path



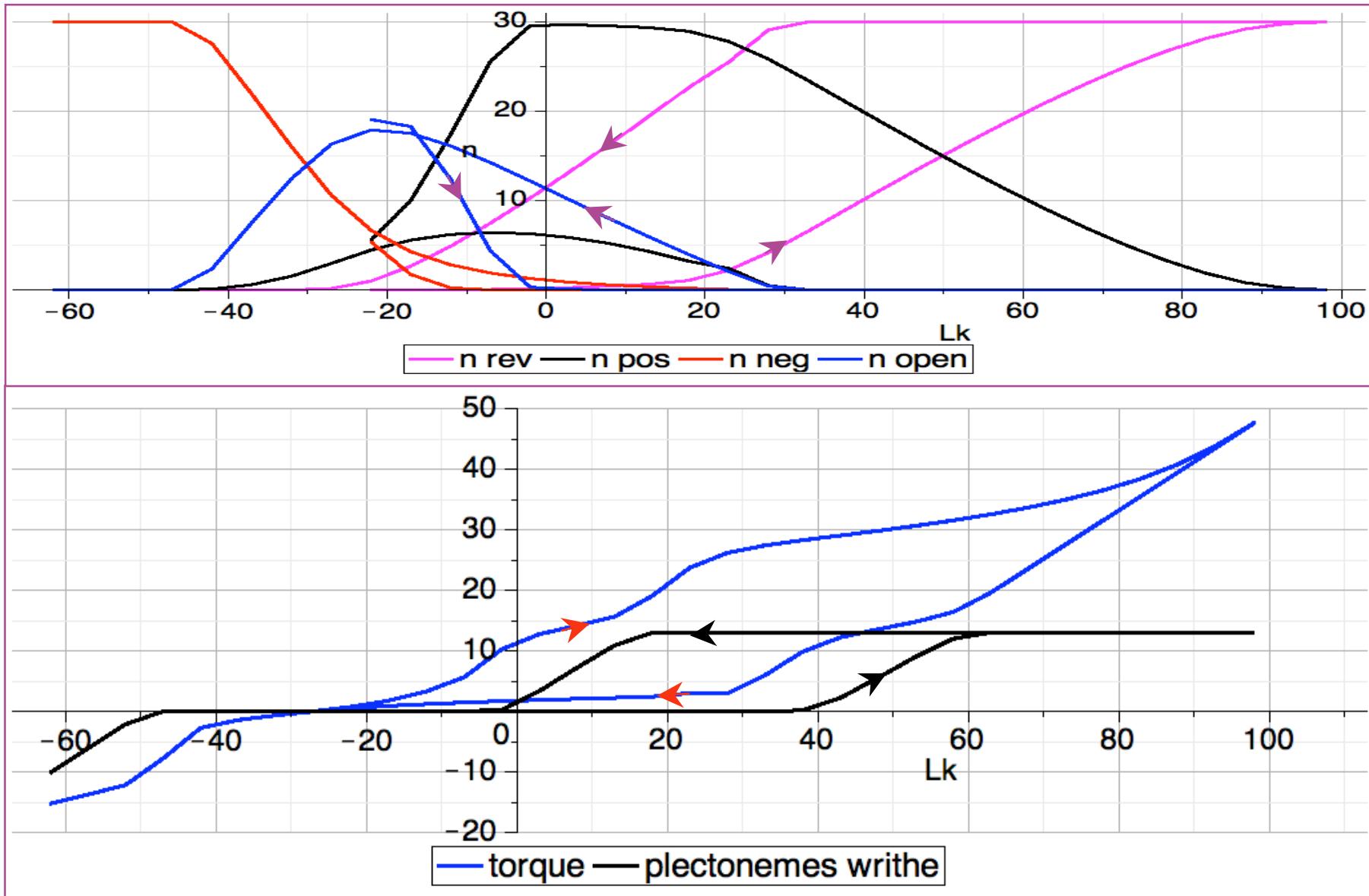
# Plectonemes formation

plectonemes initiated thanks to fiber defects via “mini-fiber” inversion:

slope:  $2 \cdot \langle L_{Minifiber} \rangle / \text{turn}$  where  $\langle L_{Minifiber} \rangle = n \cdot \langle d \rangle / (N - n + 1)$

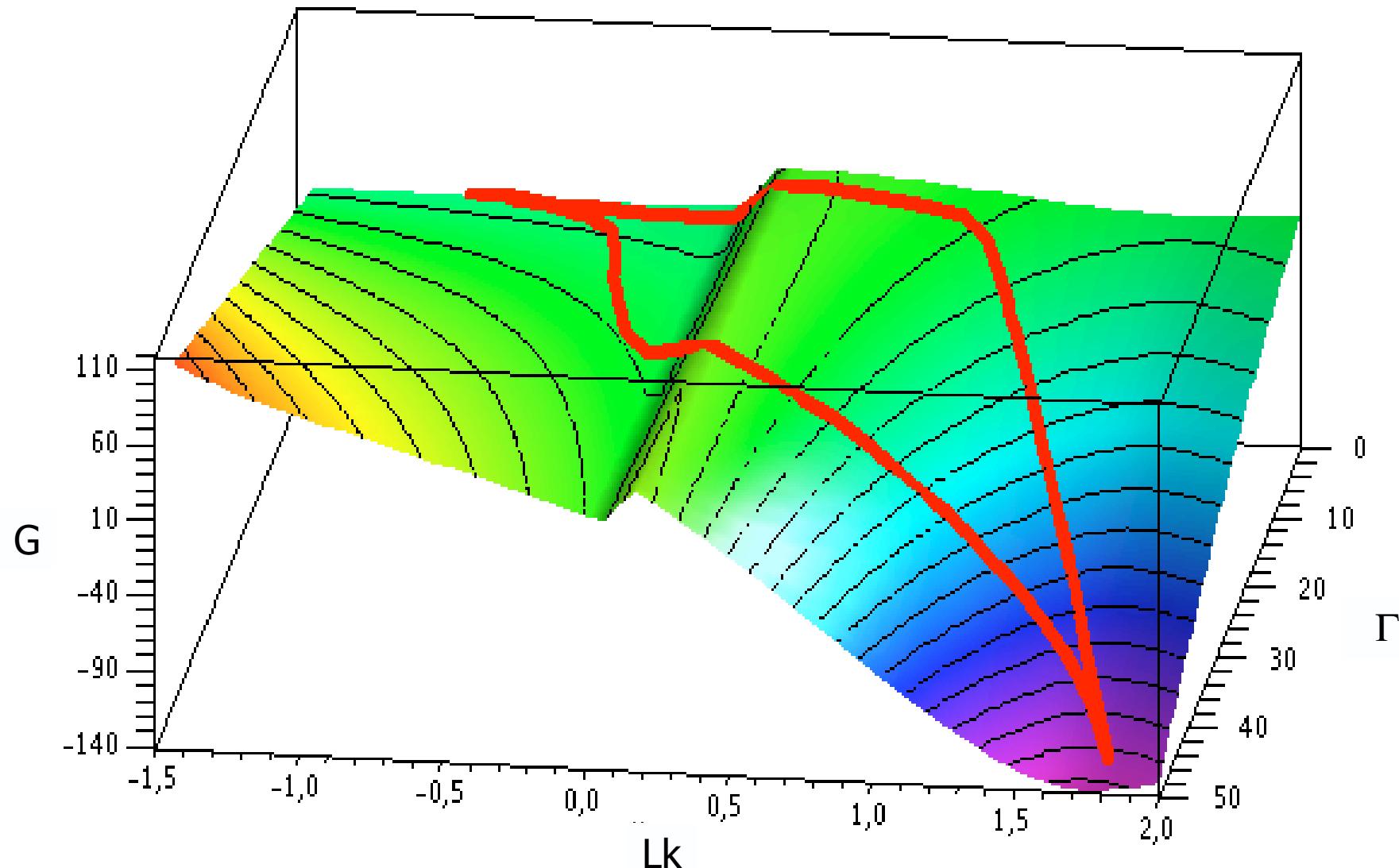


## measuring the torque via the fit

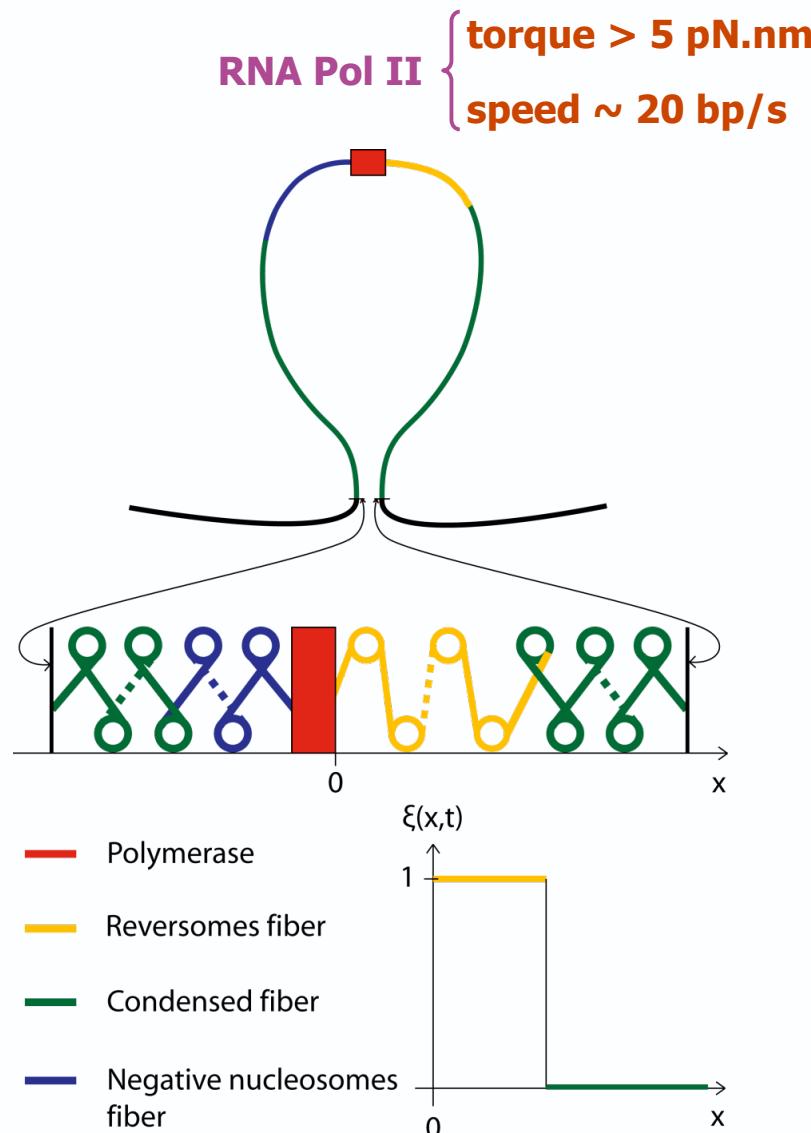


the applied torque is a model parameter which depends on  $L_k$

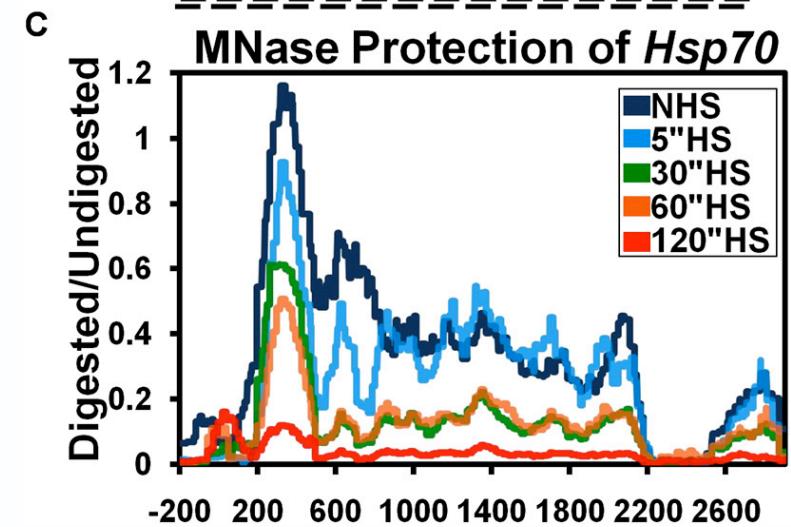
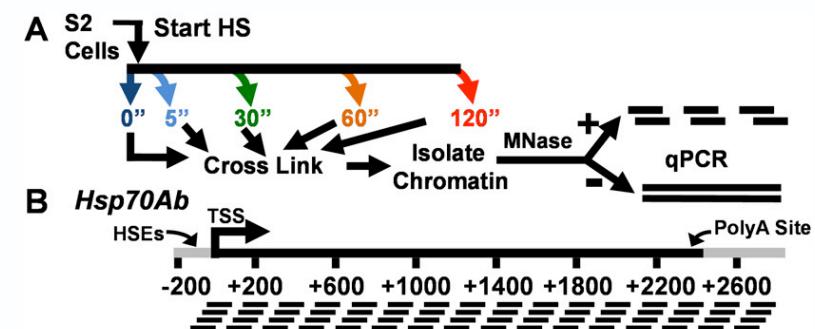
# The hysteresis cycle on the energetic landscape



# A reversome wave?



$\Rightarrow \sim 2 \text{ turns/s} \Rightarrow \sim 1 \text{ reversome/s} \Rightarrow \sim 200 \text{ bp/s}$



**A wave of nucleosome disruption**  
**(Petesh and Lis, Cell 2008)**