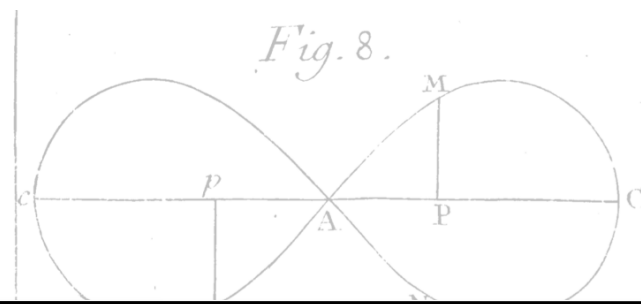
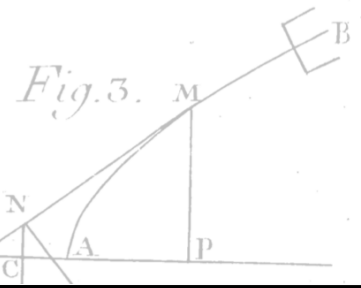
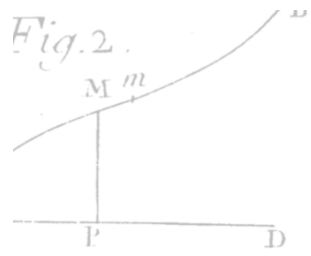


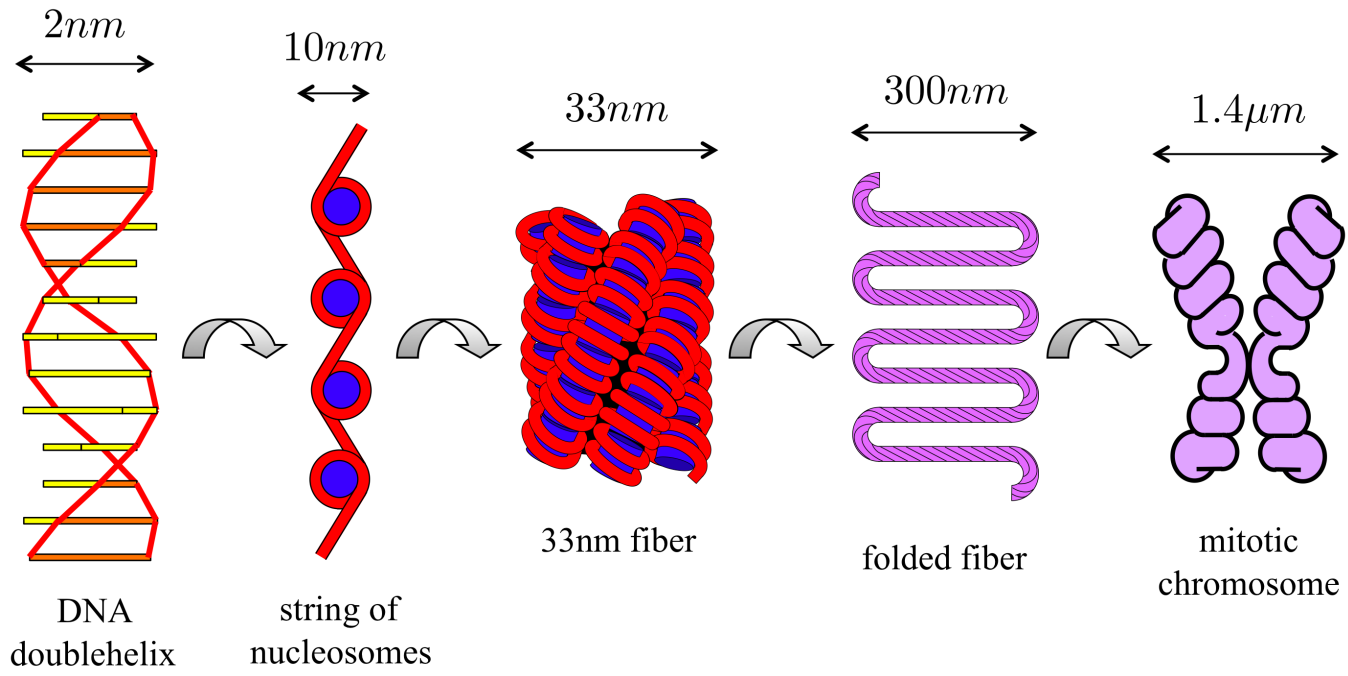
Euler's elastica and their application to DNA, nucleosomes and chromatin fibers

Helmut Schiessel

Universiteit Leiden
 Instituut Lorentz
 The Netherlands



Overview

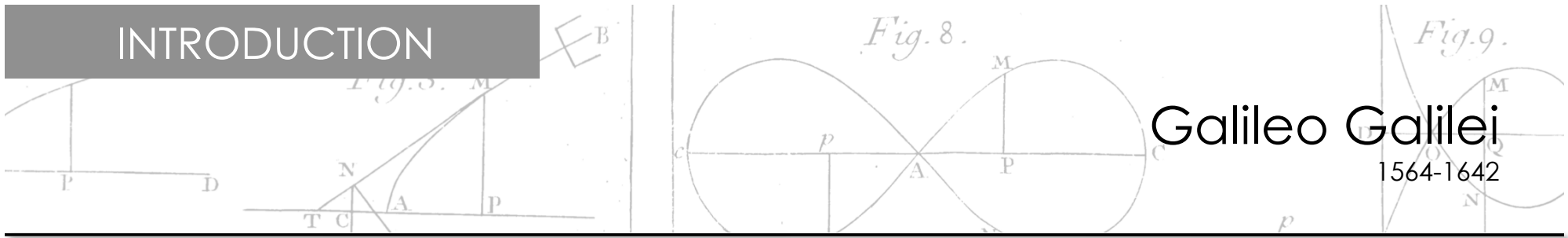


1

2

3

INTRODUCTION



Galileo Galilei
1564-1642

Discorsi, 1638:



INTRODUCTION

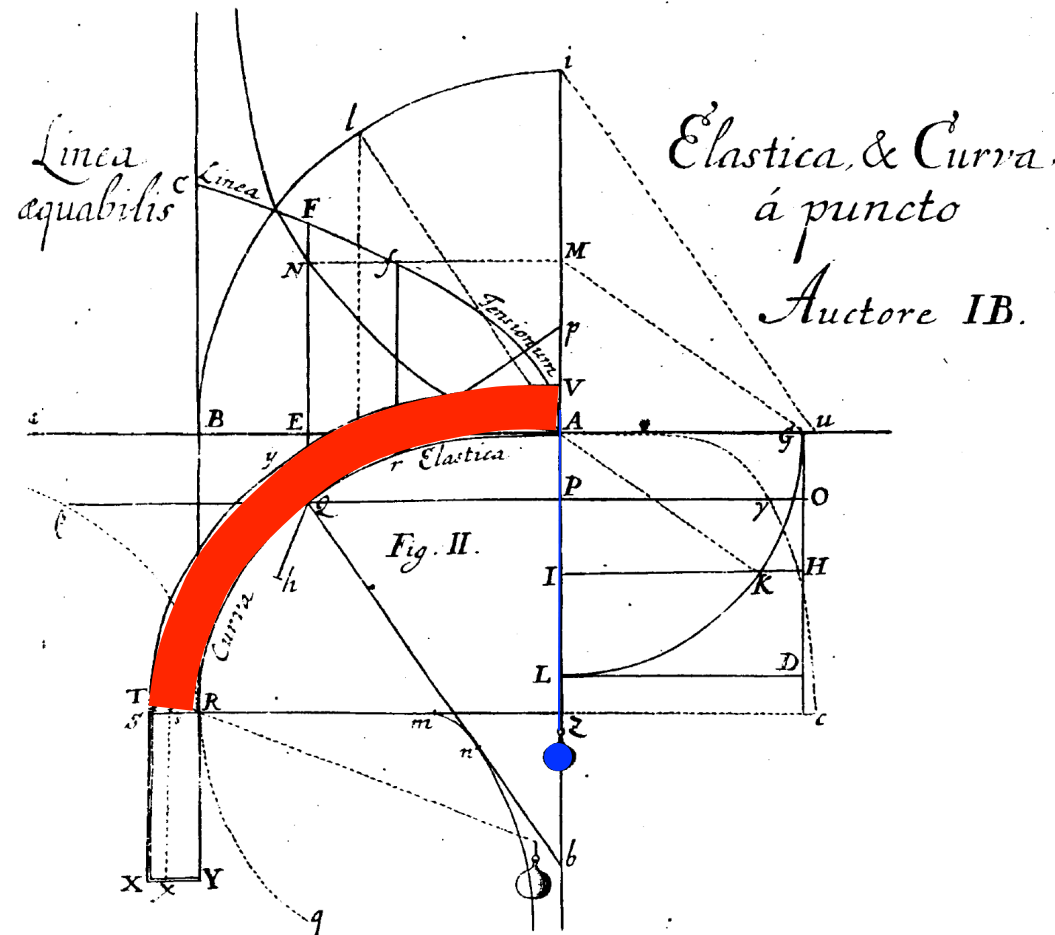
Fig. 8.

Fig. 9.

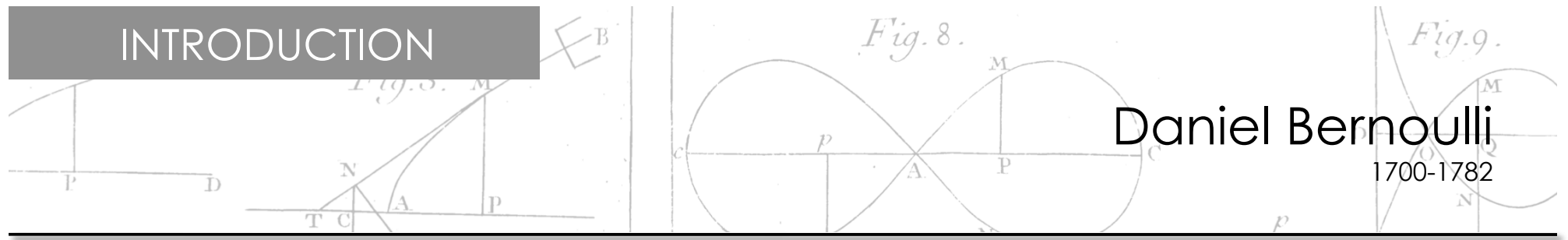
Jacob Bernoulli

1654-1705

Curvatura Laminae Elasticae, 1694:



INTRODUCTION



Daniel Bernoulli

1700-1782

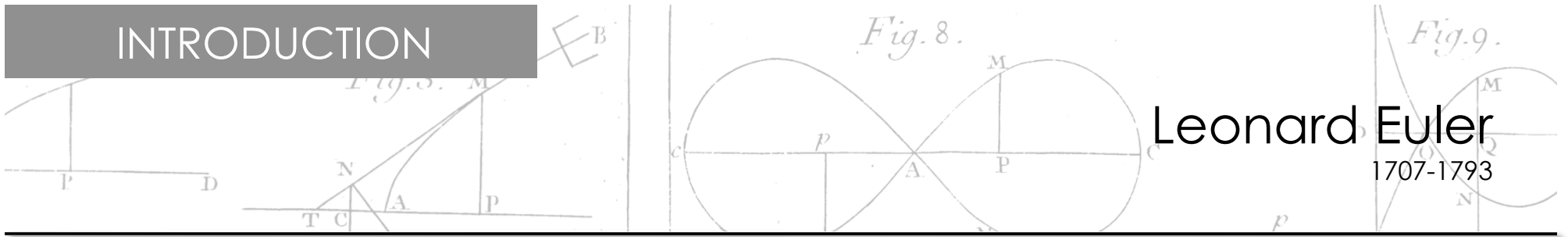
Letter to Euler, October 1742:

I'd like to know whether you might not solve the curvature of the elastic lamina under this condition, that on the length of the lamina on two points the position is fixed, and that the tangents at these points are given.

[...]

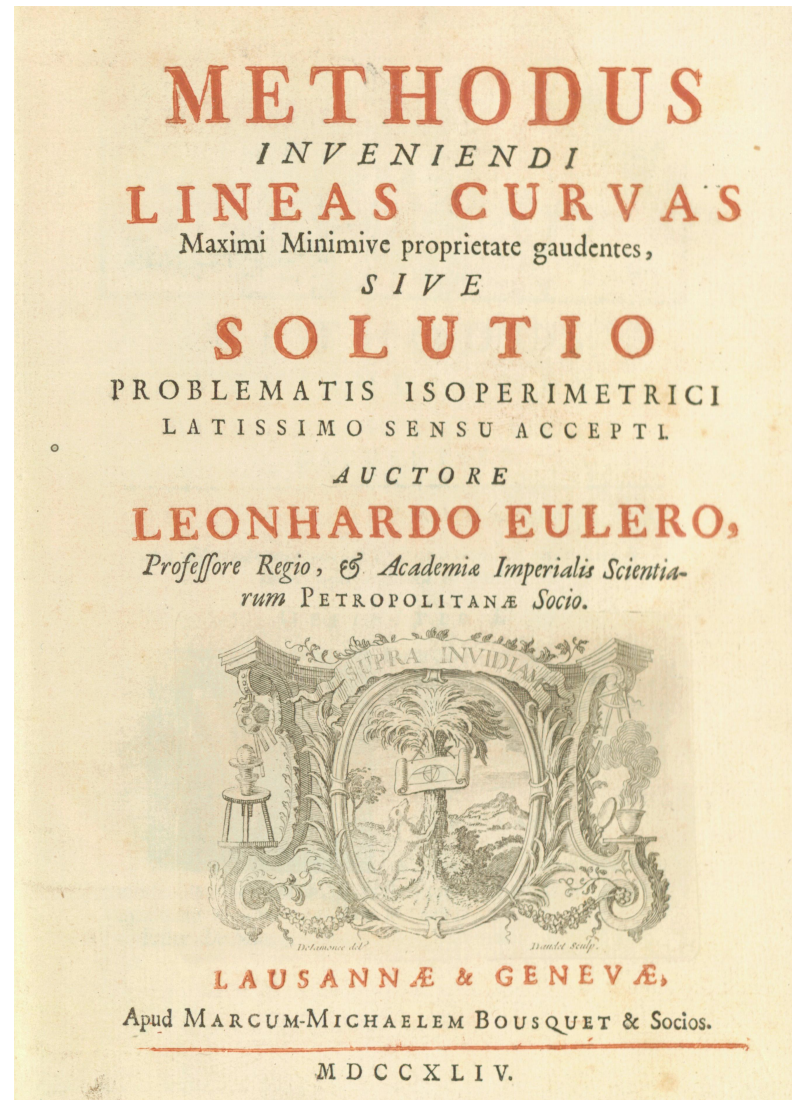
I'd express the potential energy of a curved elastic lamina (which is straight when in its natural position) through $\int \frac{ds}{RR}$, assuming the element ds is constant and indicating the radius of curvature by R . There is nobody as perfect as you for easily solving the problem....

INTRODUCTION



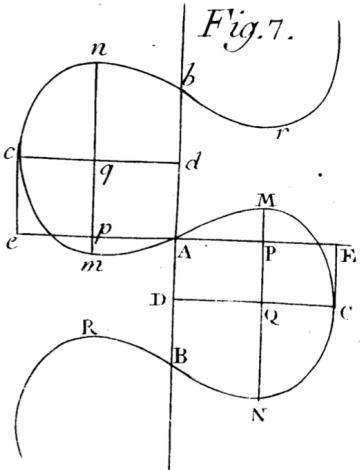
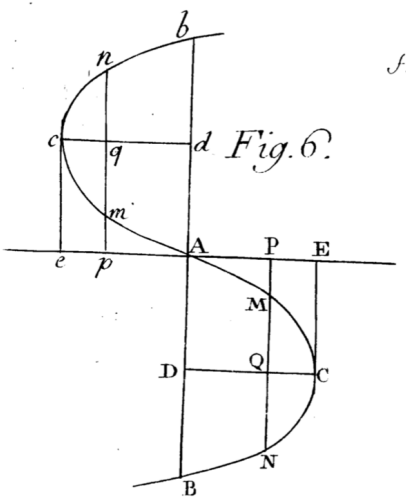
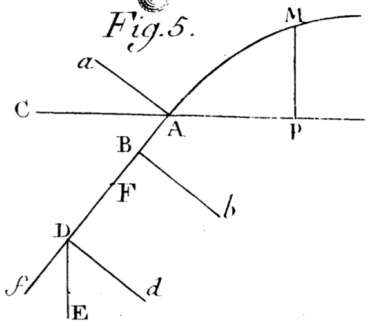
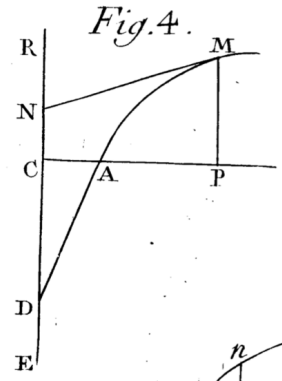
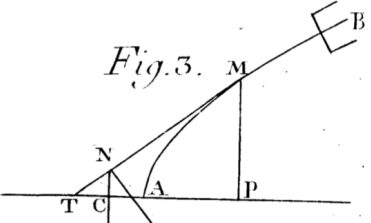
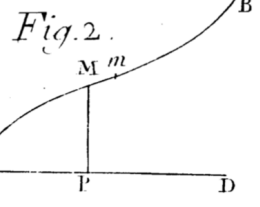
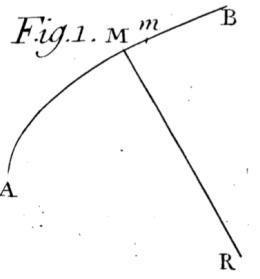
Leonard Euler
1707-1793

Euler, 1744:



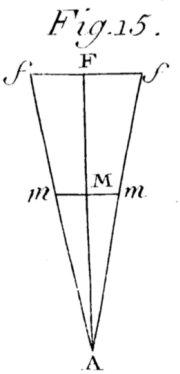
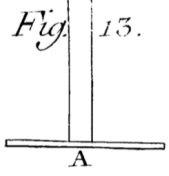
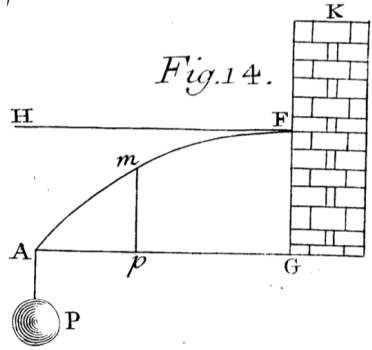
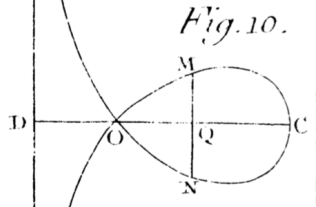
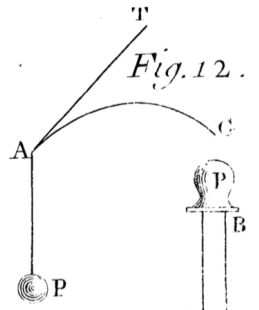
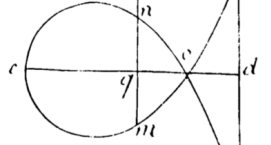
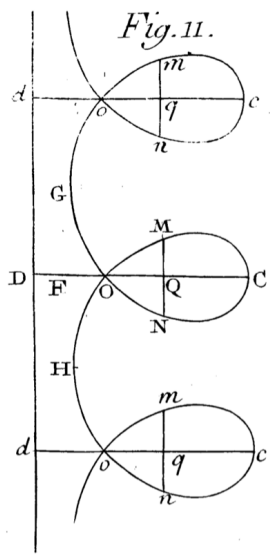
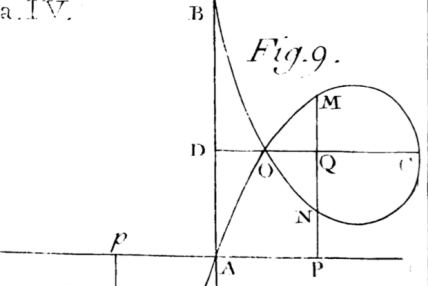
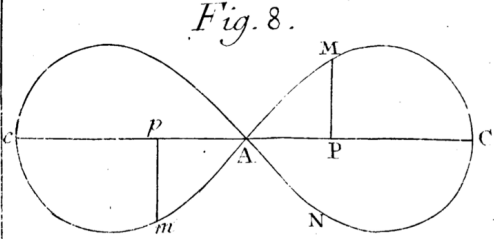
Tabula.III.

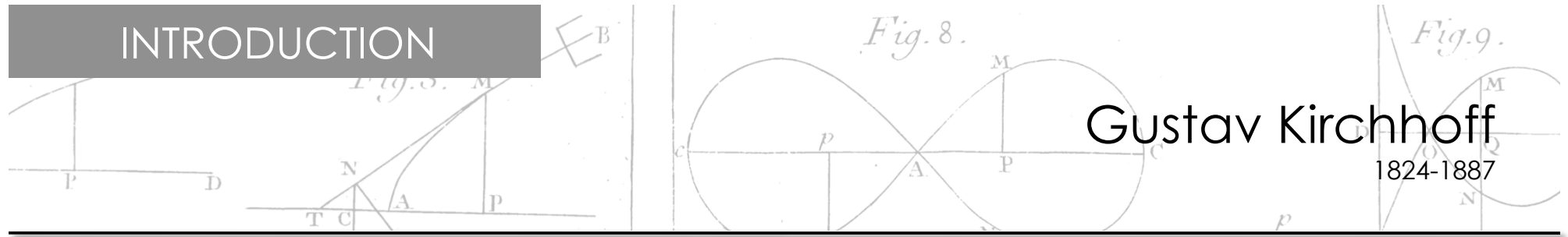
Additamentum.



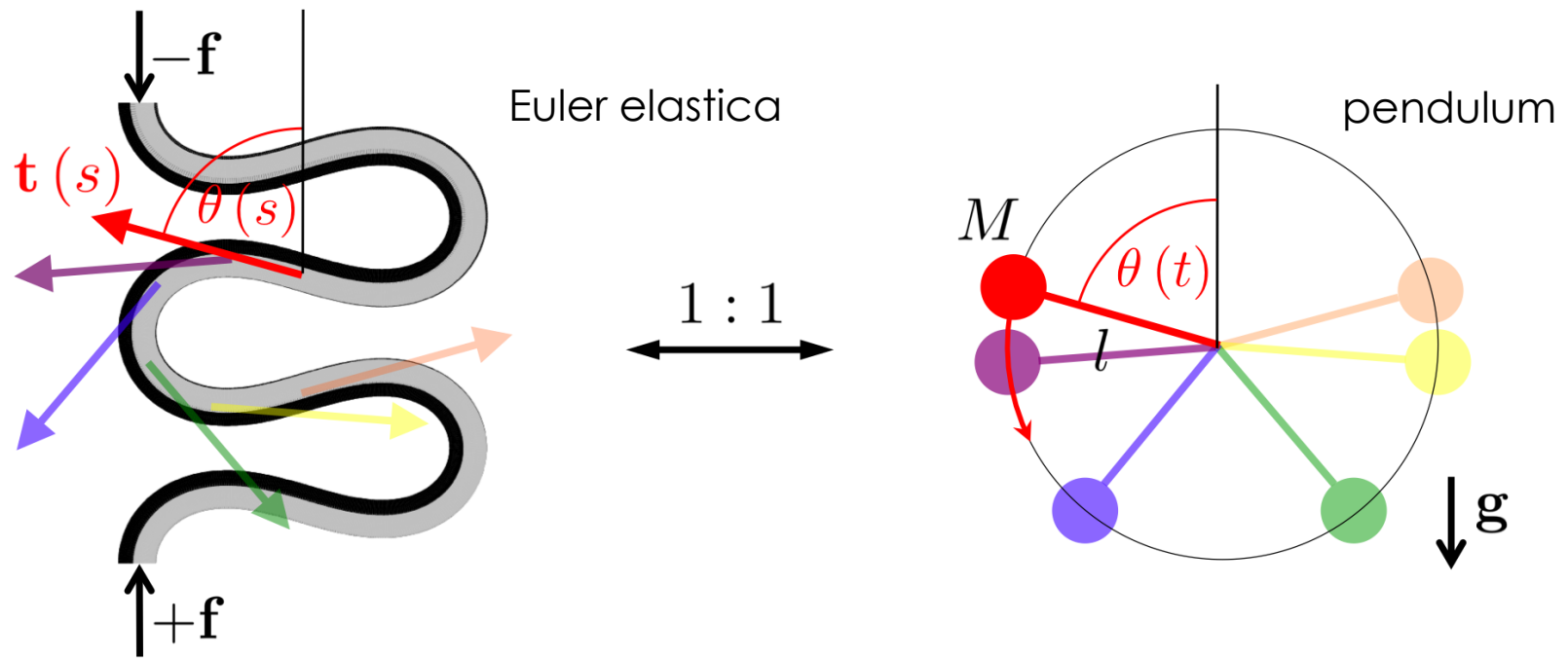
Tabula.IV.

Additamentum.





Kirchhoff kinetic analogy (1859):



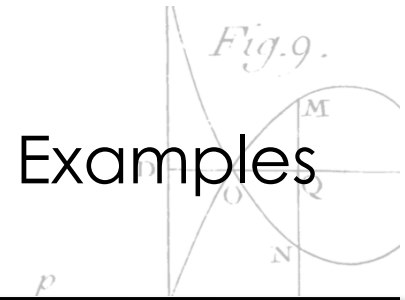
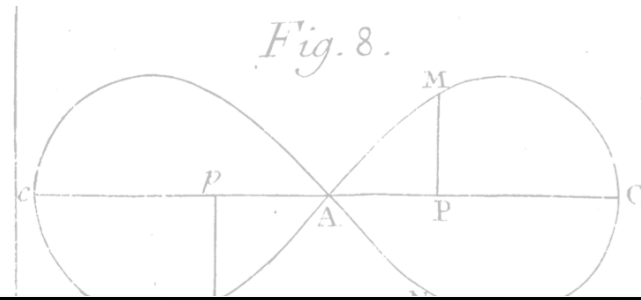
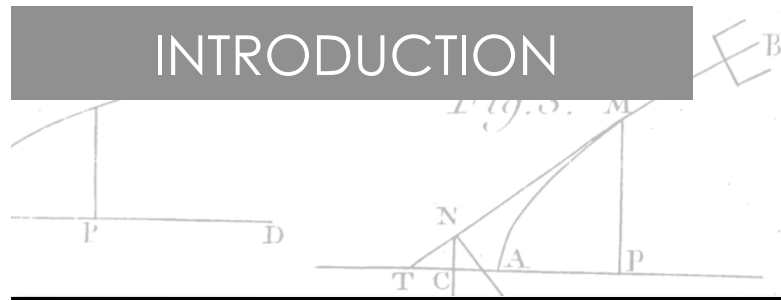
Hamiltonian (wormlike chain model):

$$H = \int_0^L ds \left[\frac{A}{2} \dot{\theta}^2 - f \cos \theta \right]$$

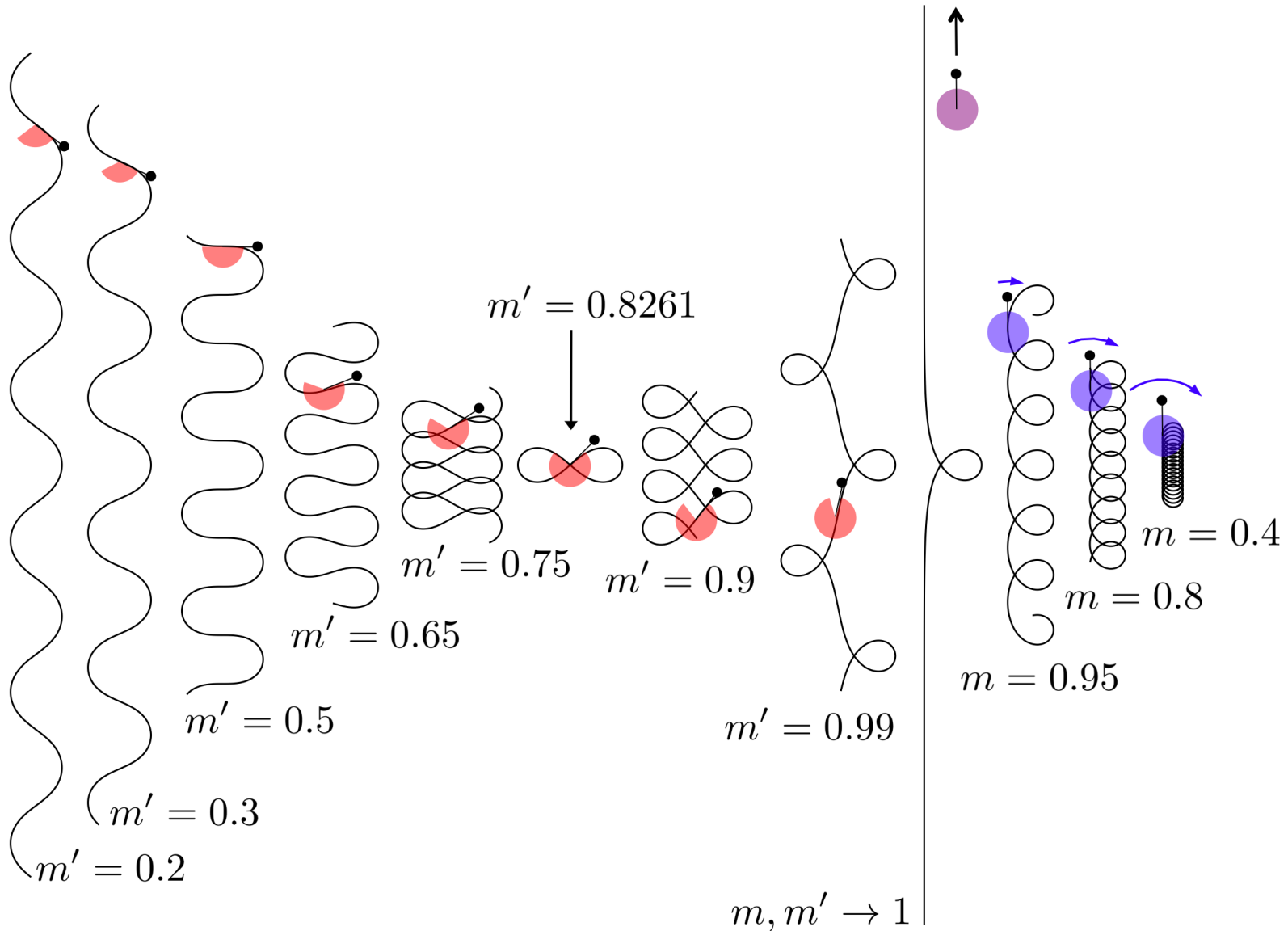
Lagrangian action:

$$S = \int_0^T d\tau \left[\frac{Ml^2}{2} \dot{\theta}^2 - Mgl \cos \theta \right]$$

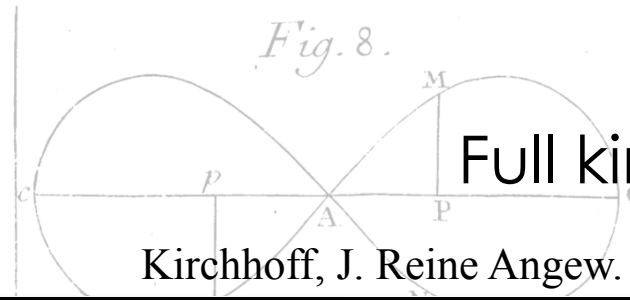
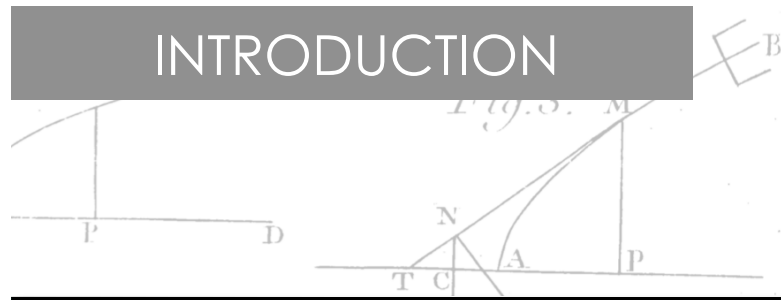
INTRODUCTION



Examples

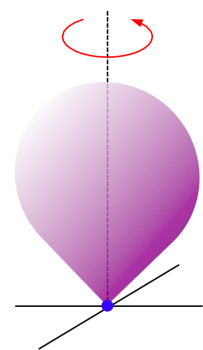


INTRODUCTION

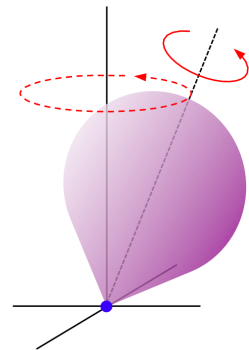


Full kinetic analogy

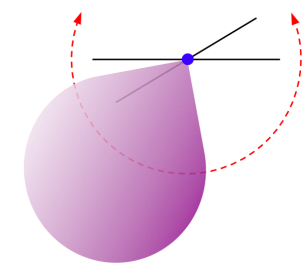
Kirchhoff, J. Reine Angew. Math **56** (1859) 285-313



sleeping top



regular precession



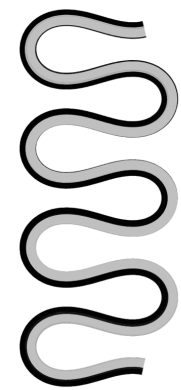
pendulum



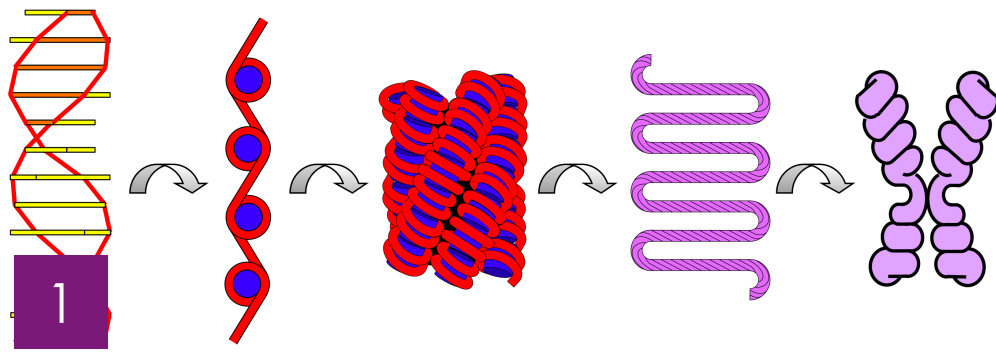
twisted rod



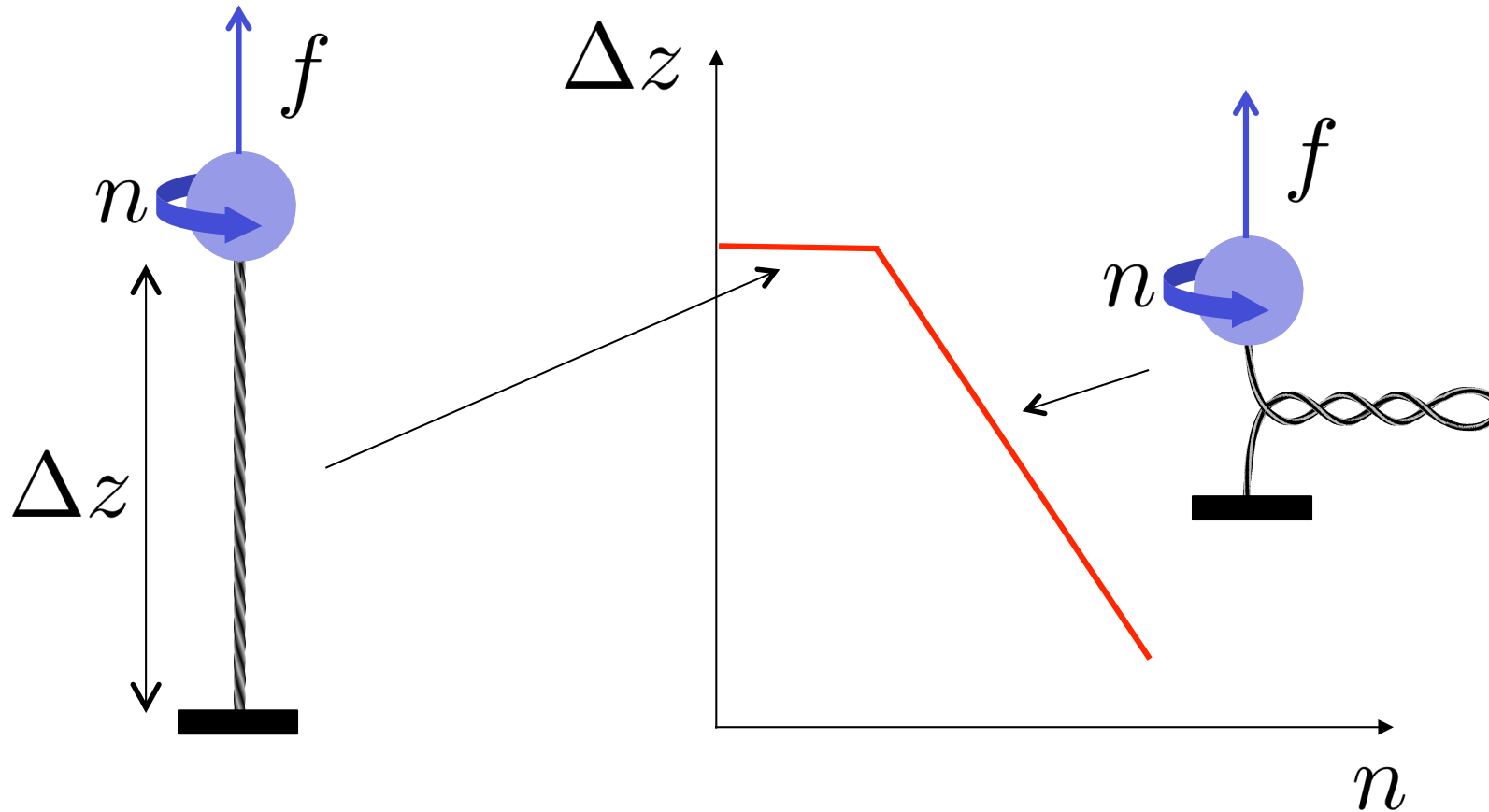
helix



planar filament



Twisting DNA under tension



Theories: Marko, Phys. Rev. E **76** (2007) 021926

Clauvelin, Audoly & Neukirch, Biophys. J. **96** (2009) 3716

Maffeo, Schopflin, Brutzer, Stehr, Aksimentiev, Wedemann & Seidel, PRL **105**, 158101

1



Fig.3.

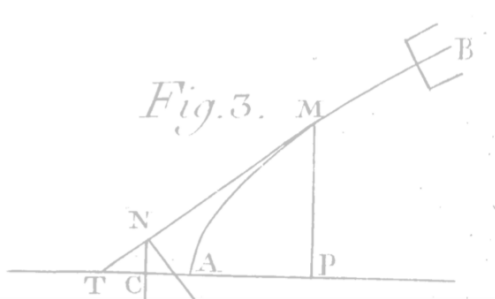


Fig.8.

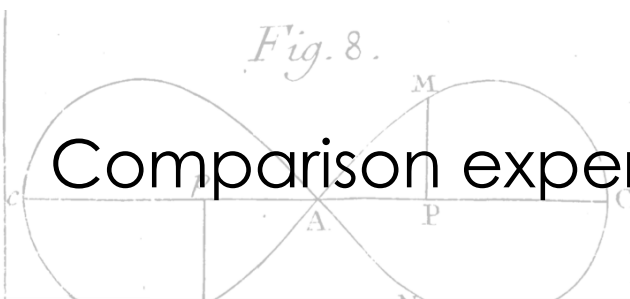
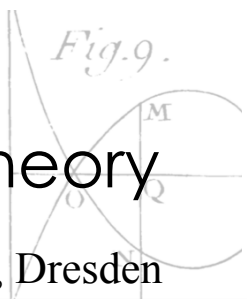


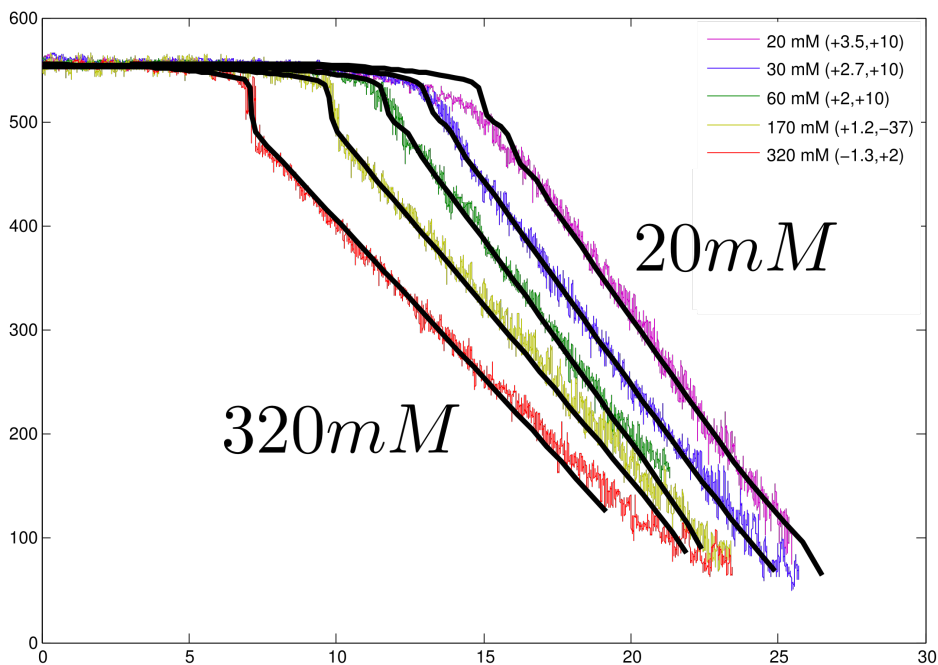
Fig.9.



Comparison experiment/theory

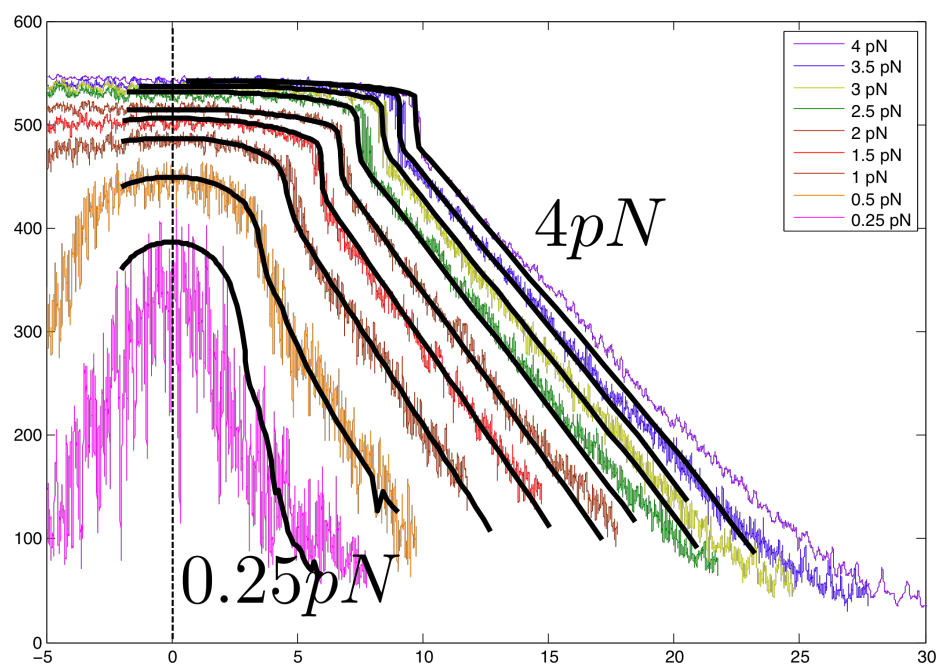
Ralf Seidel lab, Dresden

Δz [nm]



$f = 3pN$

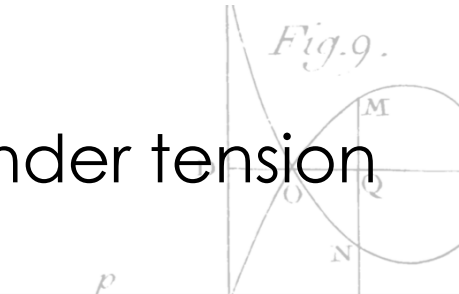
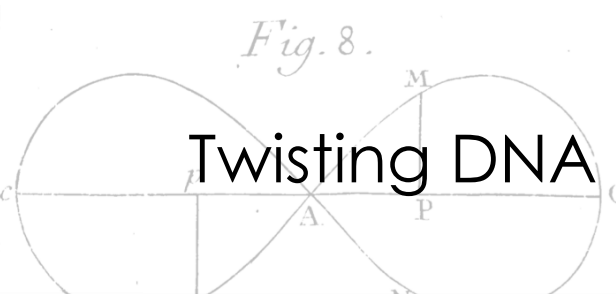
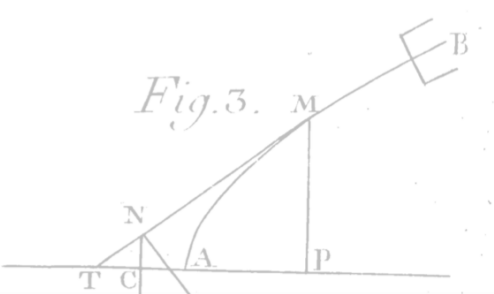
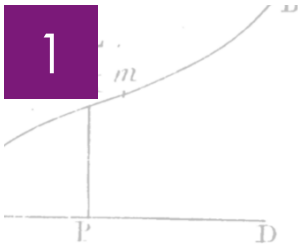
Δz [nm]



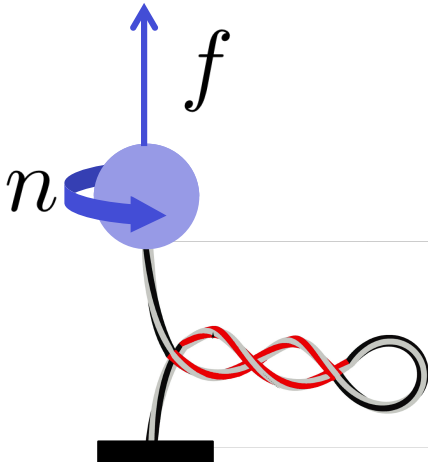
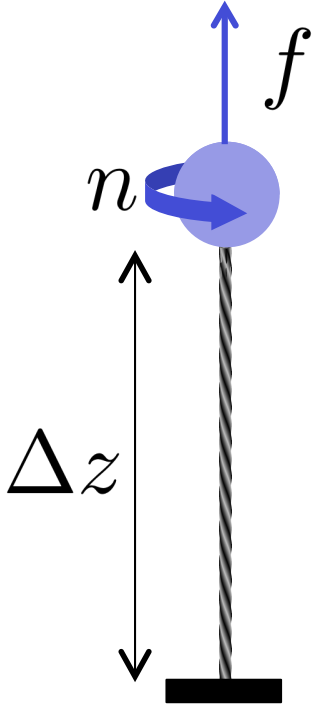
$c_s = 320mM$

$L = 600nm$

1



Twisting DNA under tension



total energy:

$$E_{\text{tot}} = \underbrace{\frac{A}{2} \int_0^L \frac{ds}{R^2(s)}}_{\text{DNA bending}} + \underbrace{\frac{C}{2} \int_0^L \Delta\psi^2(s) ds}_{\text{DNA twisting}} - \underbrace{f \Delta z}_{\text{external force}} - \underbrace{2\pi n ([\mathbf{t}, \Delta\psi]) \tau_f}_{\text{external twist}} + \underbrace{U_{\text{el}}([\mathbf{t}])}_{\text{electrostatic contribution}}$$

1

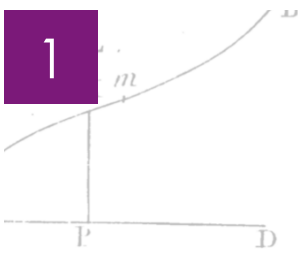


Fig. 3.

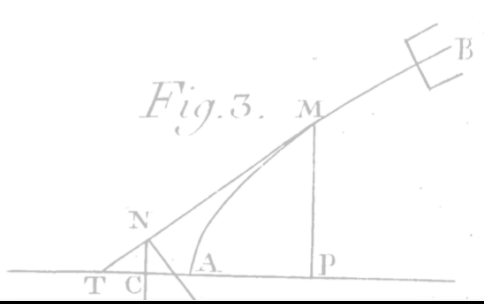


Fig. 8.

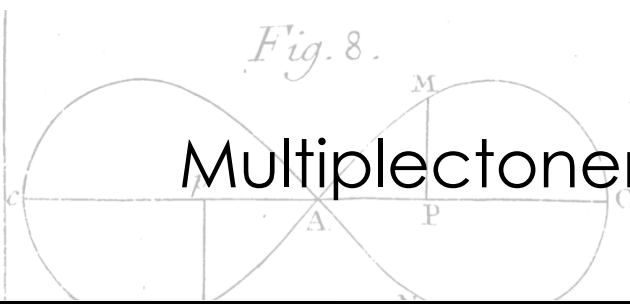
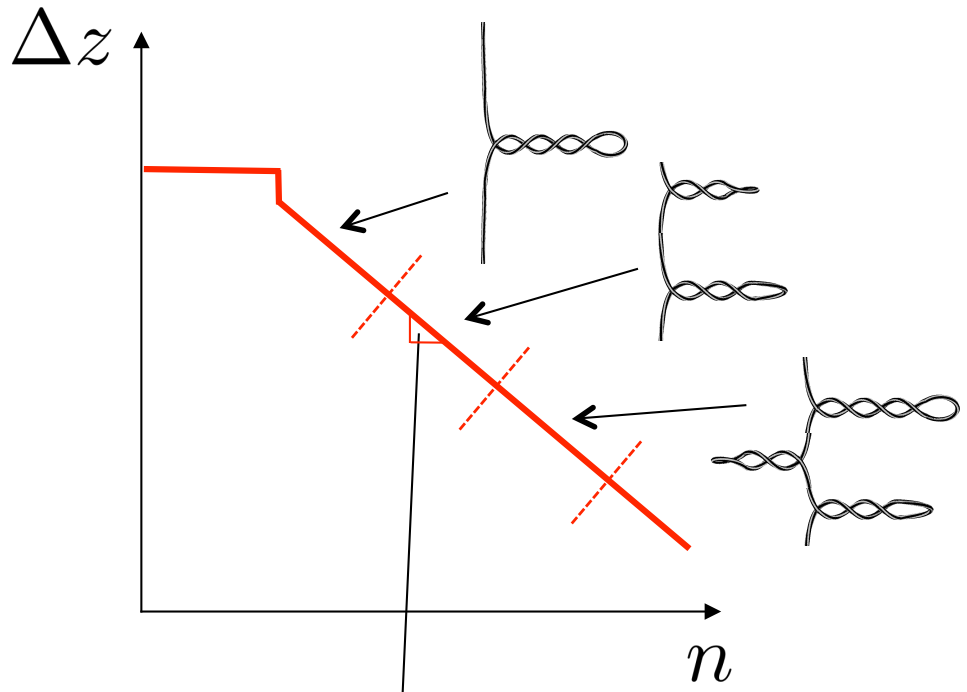
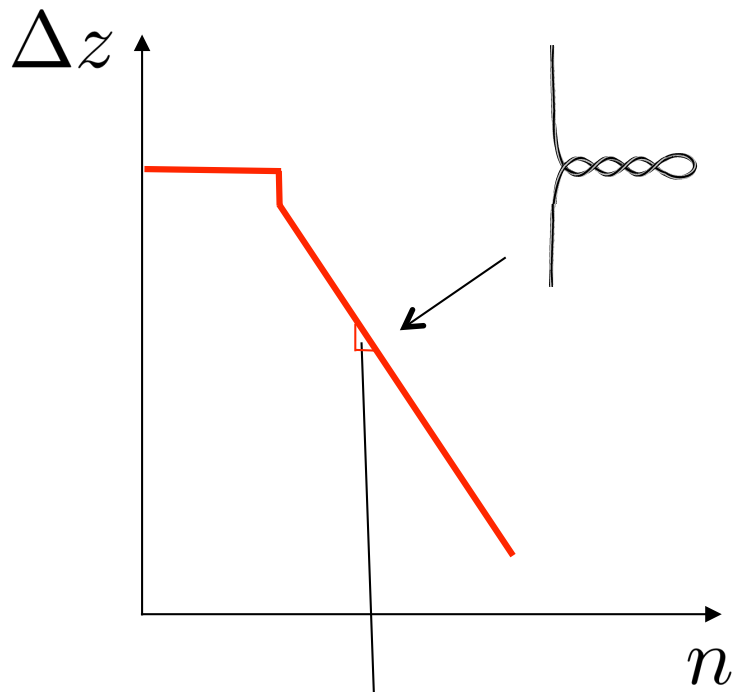


Fig. 9.

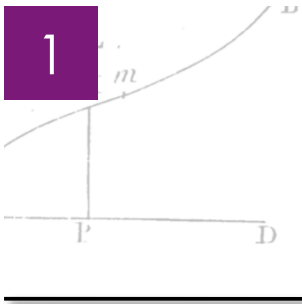
Multiplectoneme formation



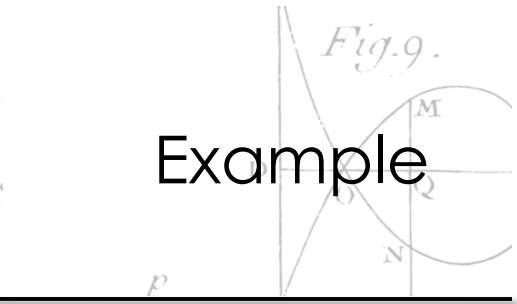
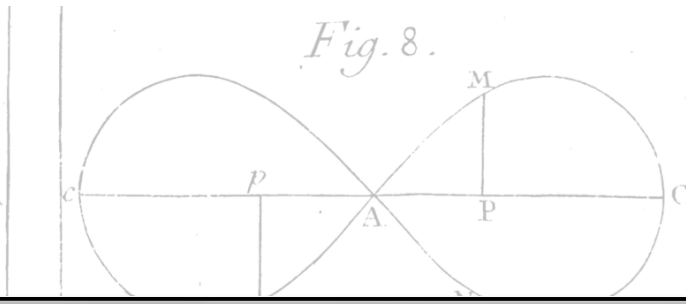
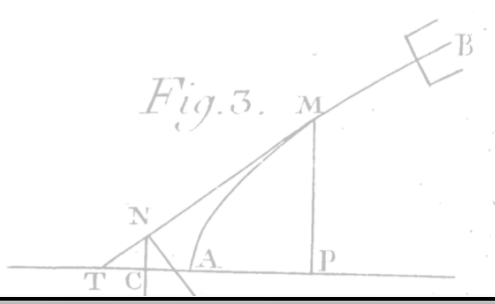
slope:

$$\frac{\partial \Delta z}{\partial n} = -\frac{\rho}{\rho_{pl}WR}$$

$$\frac{\partial \Delta z}{\partial n} = -\frac{\rho}{\rho_{pl}WR} + \frac{\partial m}{\partial n} \left(\frac{\rho}{\rho_{pl}WR} - l_{loop} \right)$$

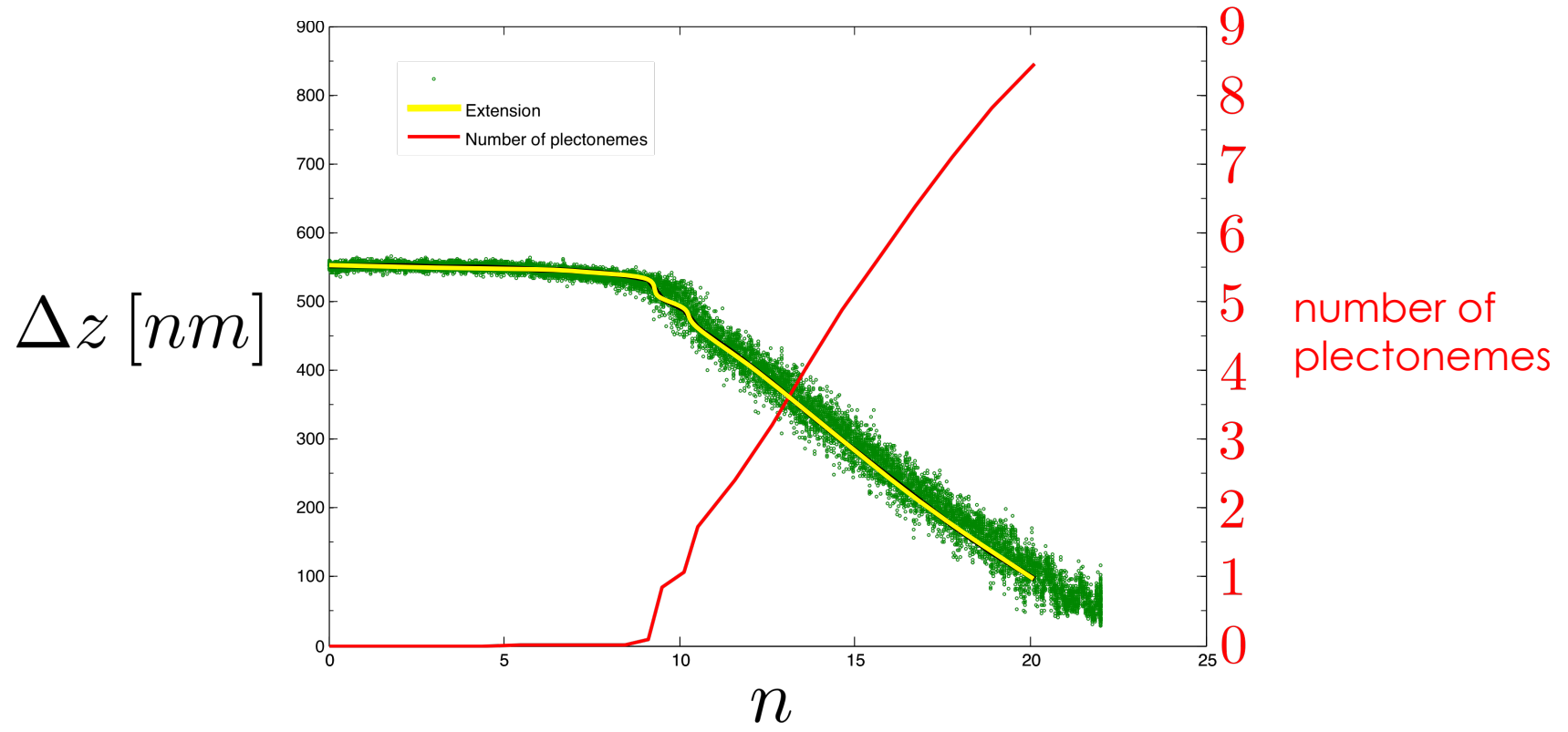


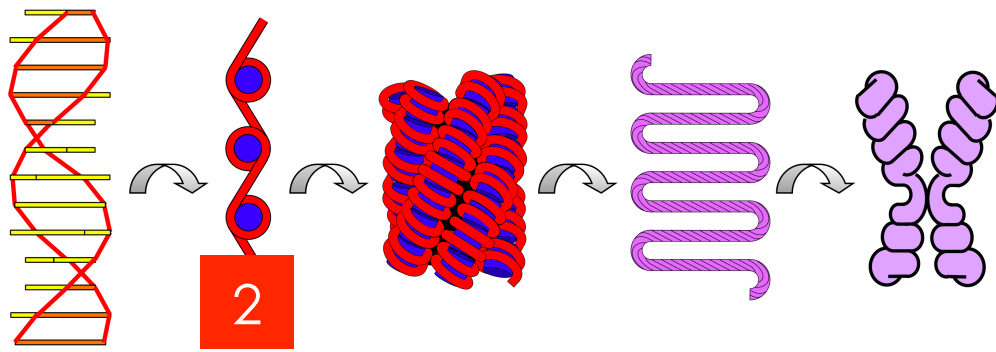
1



Example

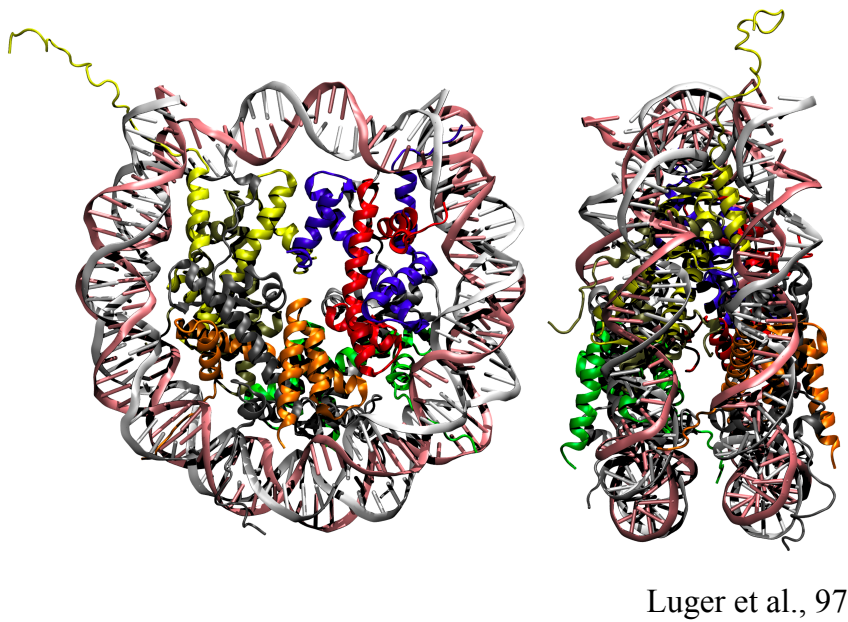
$$L = 600nm \quad f = 3pN \quad c_s = 30mM$$



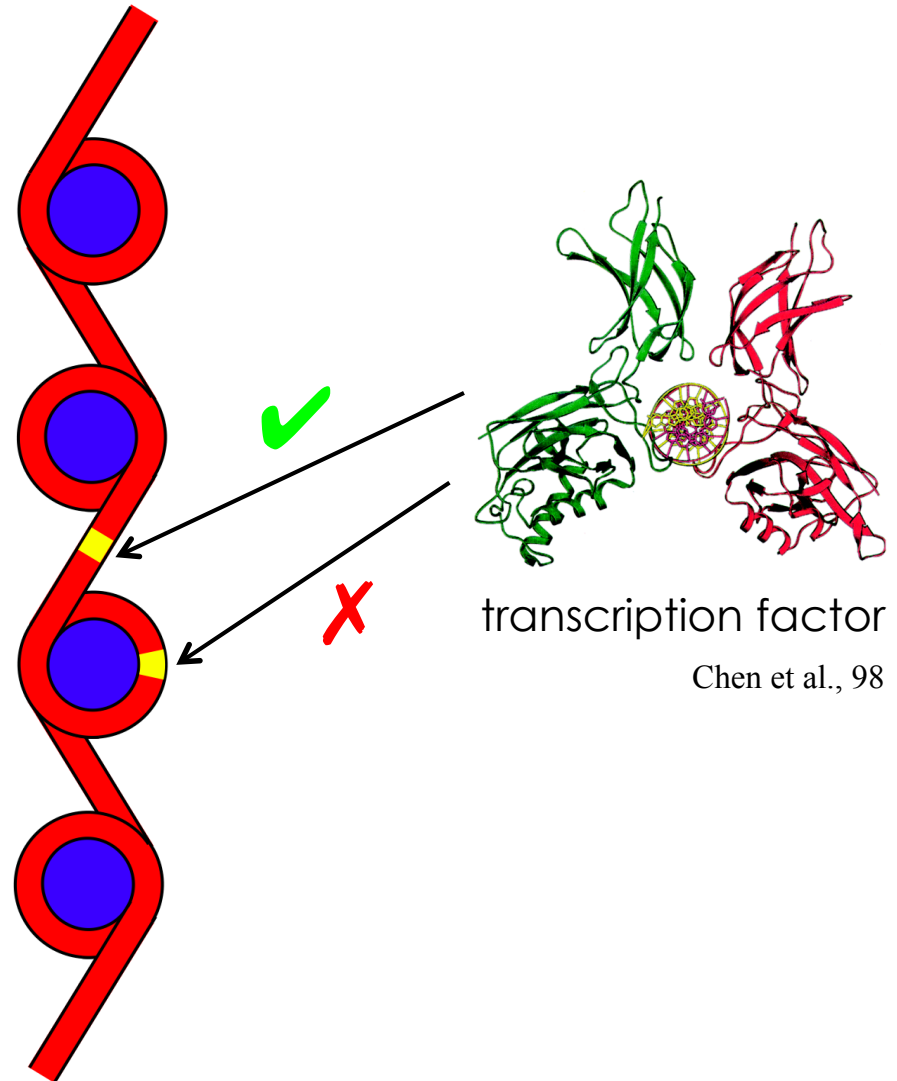


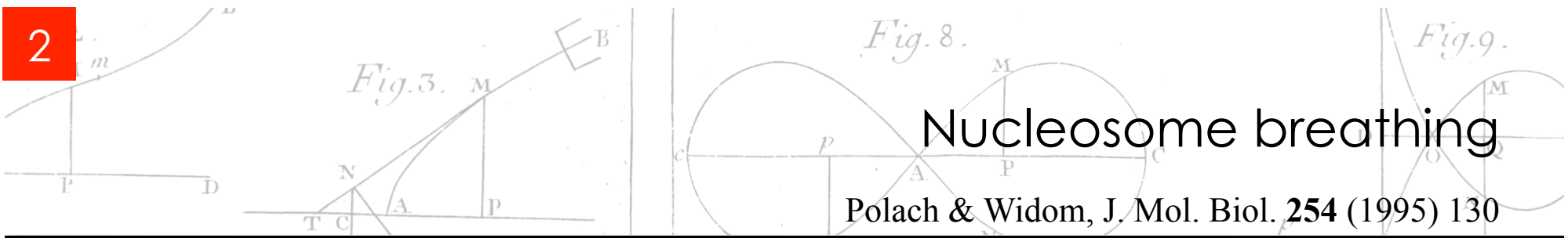
Nucleosomes under tension

nucleosome core particle:



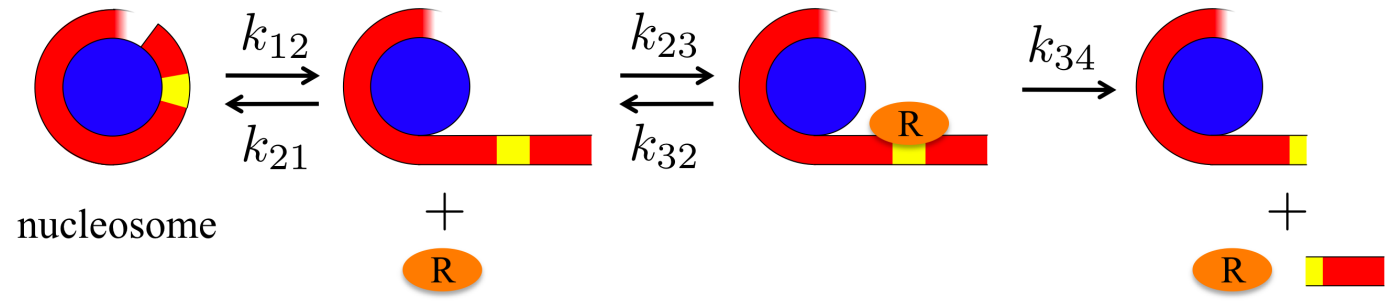
- 1 3/4 turns
- 147 bp, 50nm
- 14 binding sites





Nucleosome breathing

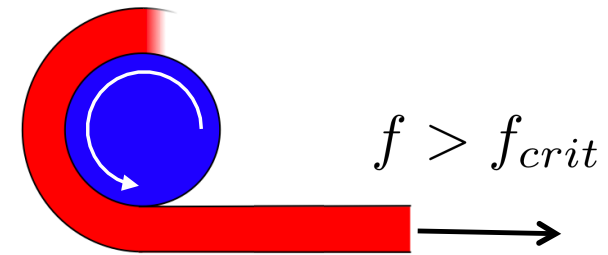
Polach & Widom, J. Mol. Biol. **254** (1995) 130

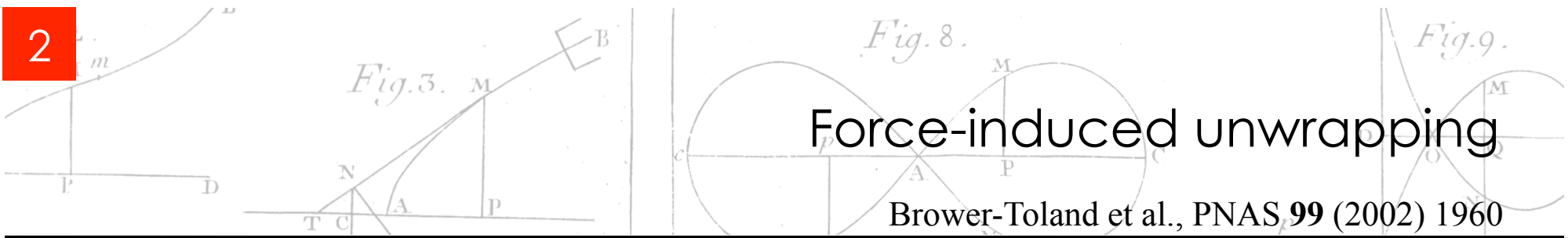


adsorption energy per length:

Prinsen & HS, Biochimie **92** (2010) 1722

$$f_{crit} \approx \frac{15k_B T}{50nm} = 1.2pN$$

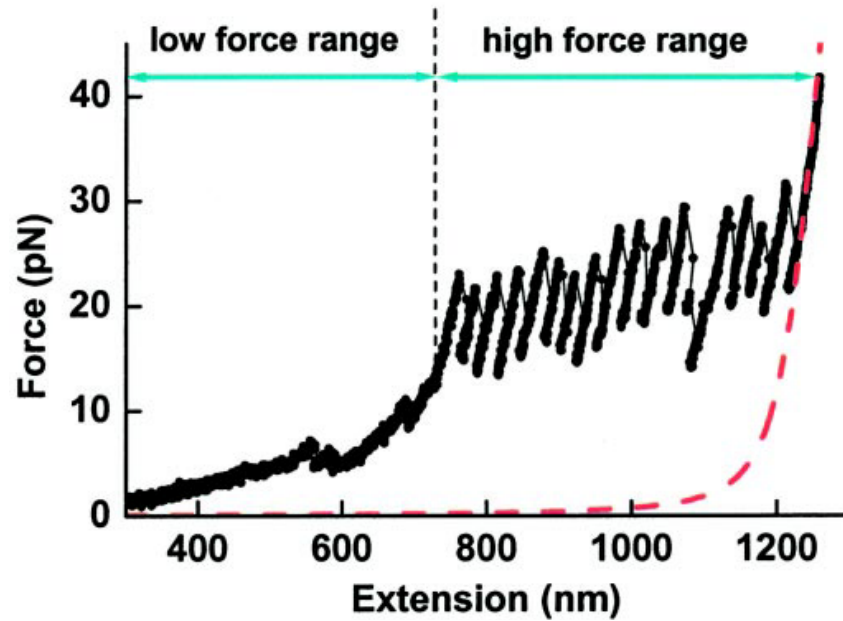
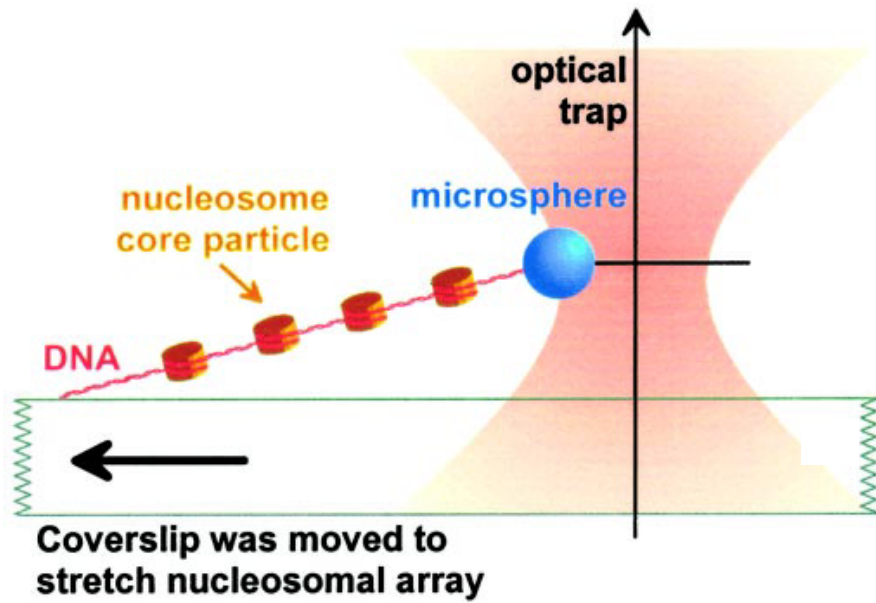




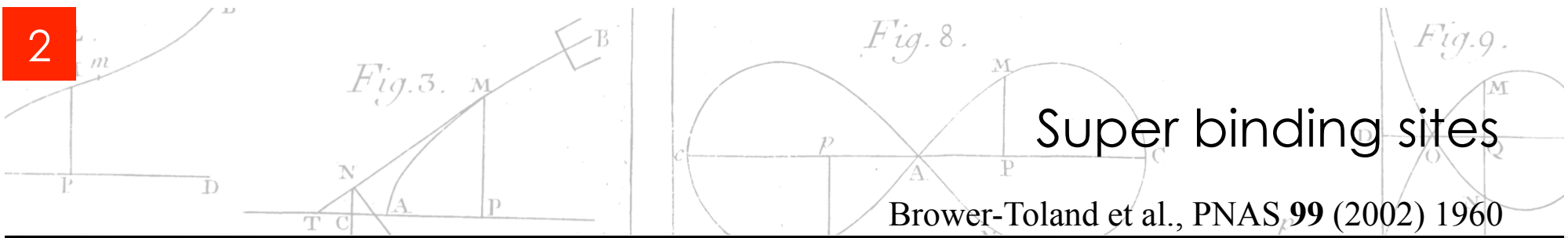
Force-induced unwrapping

Brower-Toland et al., PNAS 99 (2002) 1960

17 nucleosomes under tension:

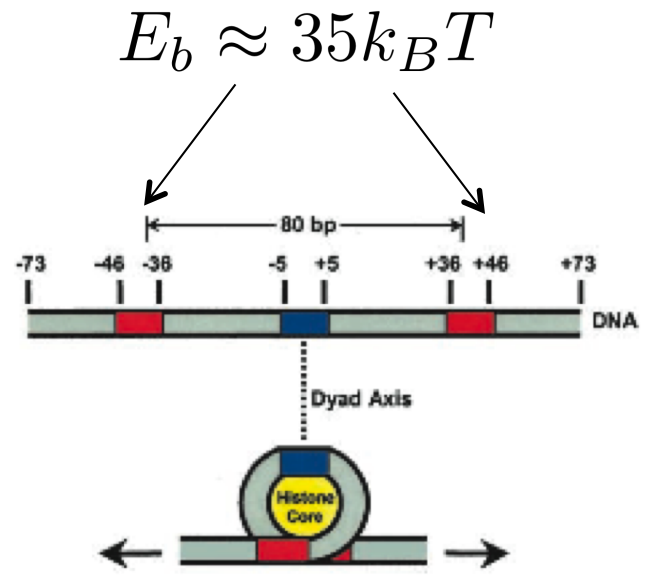
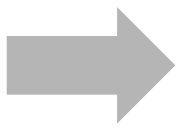
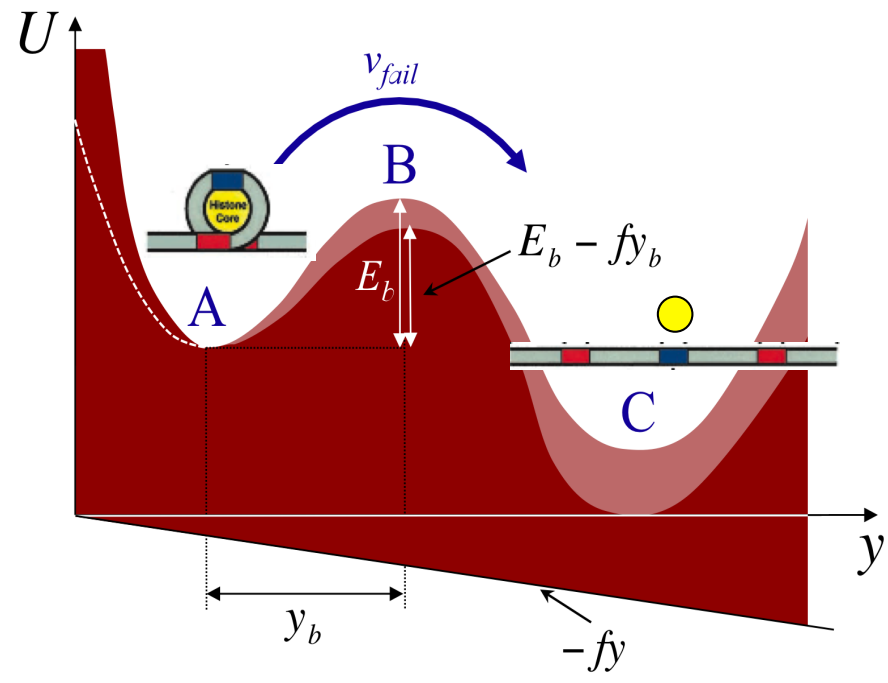


But: $f_{\text{crit}} \approx \frac{15k_B T}{50\text{nm}} = 1.2\text{pN}$

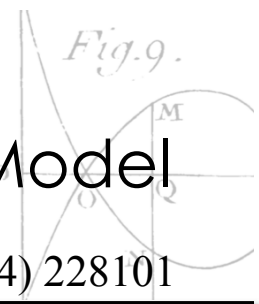
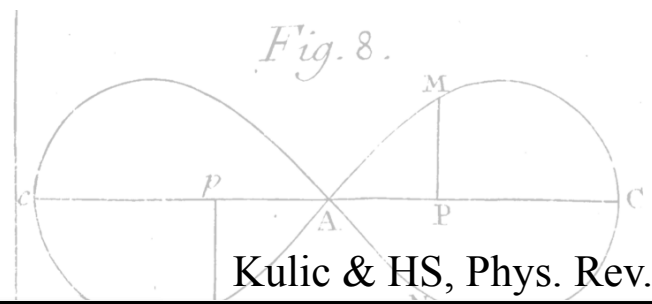
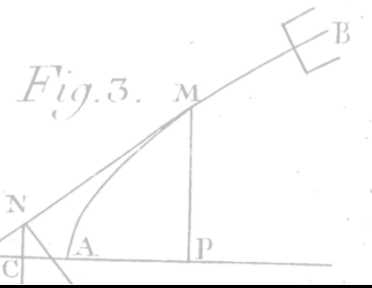
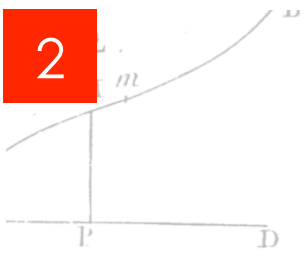


Super binding sites

Brower-Toland et al., PNAS 99 (2002) 1960

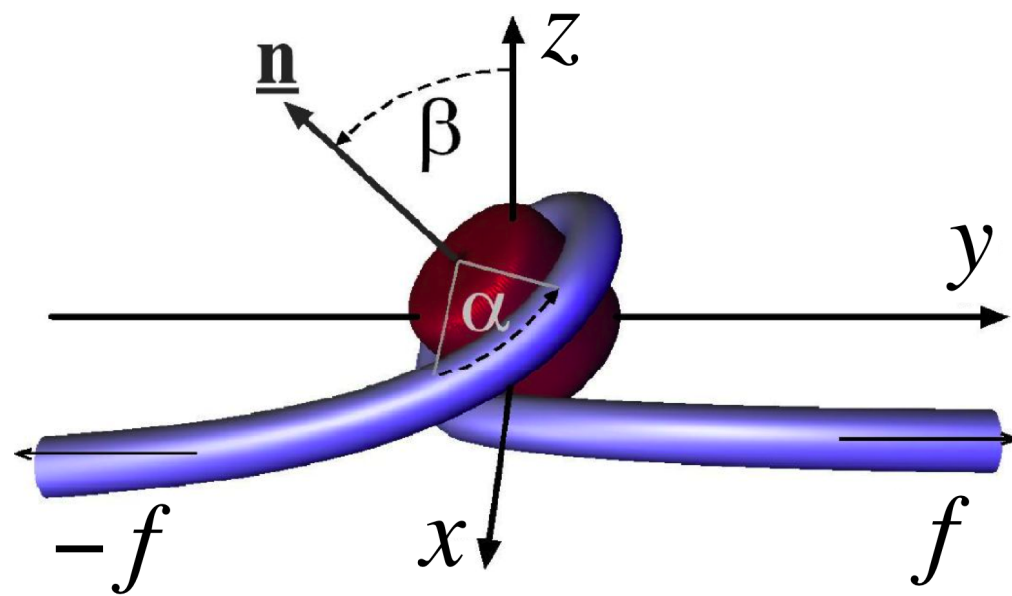


But: $E_{complex} \approx 15k_B T$

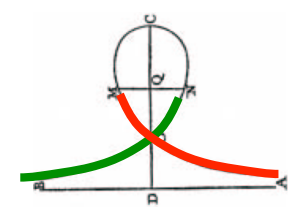


Model

Kulic & HS, Phys. Rev. Lett. **92** (2004) 228101



α : unwrapping angle
 β : tilting angle



total energy:

$$E_{\text{tot}} = \underbrace{\frac{A}{2} \int_0^L ds \left(\frac{1}{R(s)} \right)^2}_{\text{DNA bending}} - \underbrace{fL_{EE}}_{\text{external force}} + \underbrace{2R_0 f_{\text{crit}} \alpha}_{\text{DNA adsorption}}$$

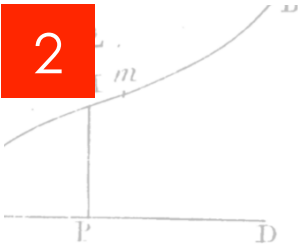


Fig. 3.

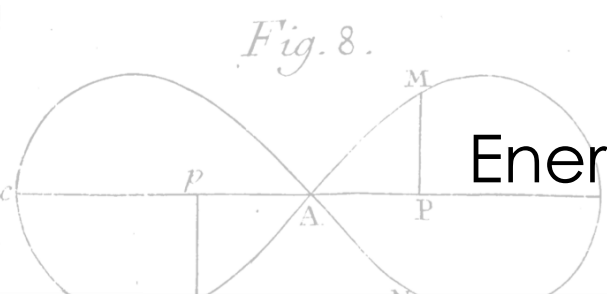


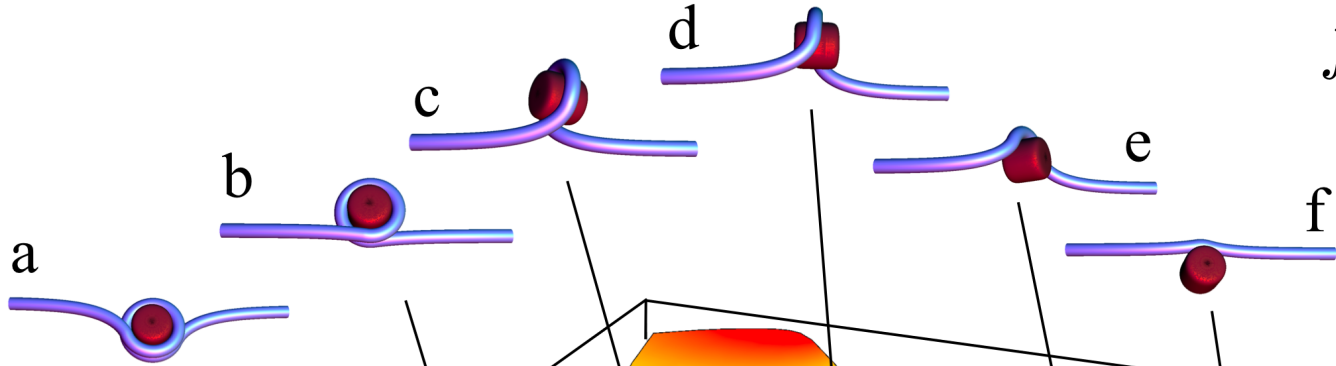
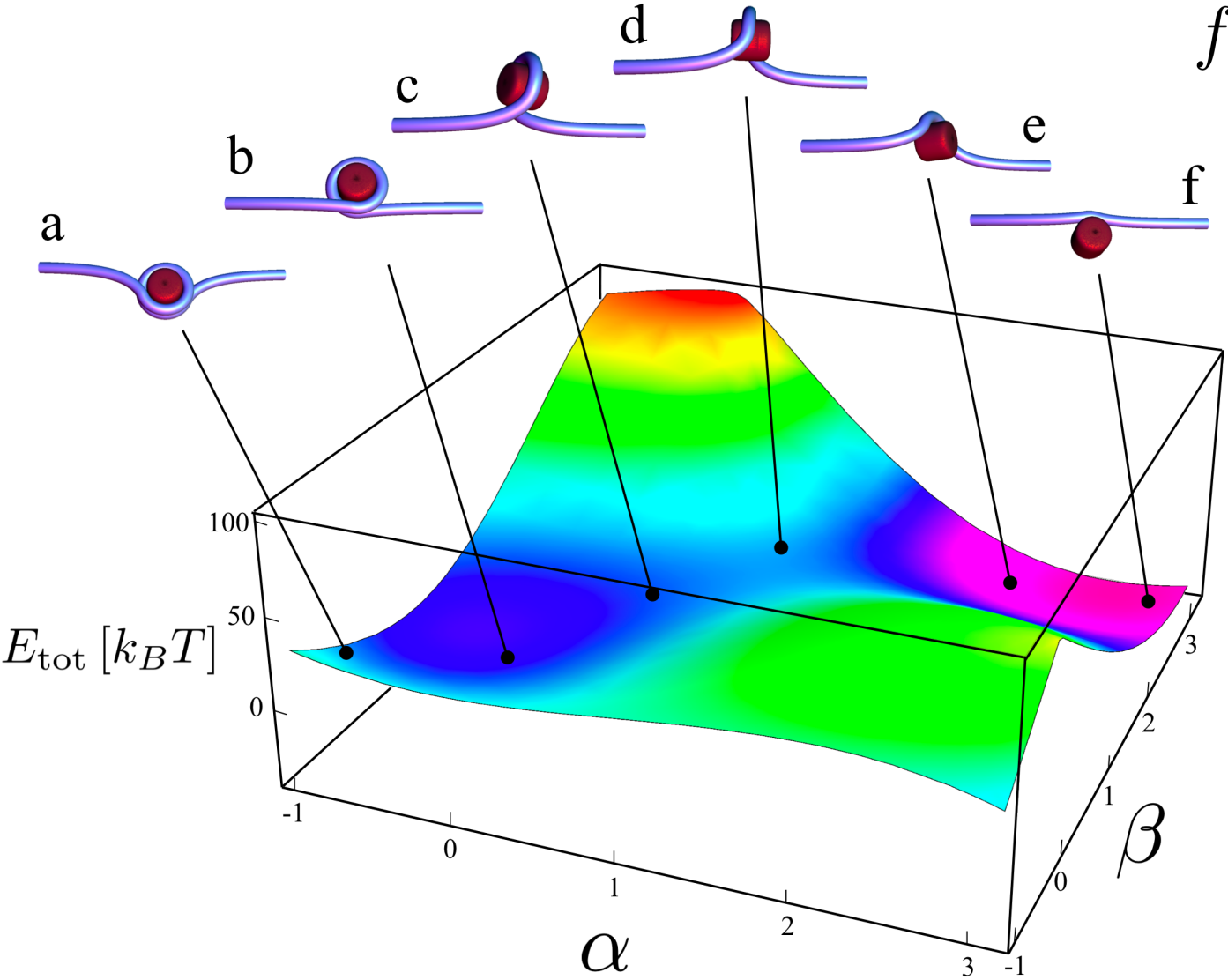
Fig. 8.

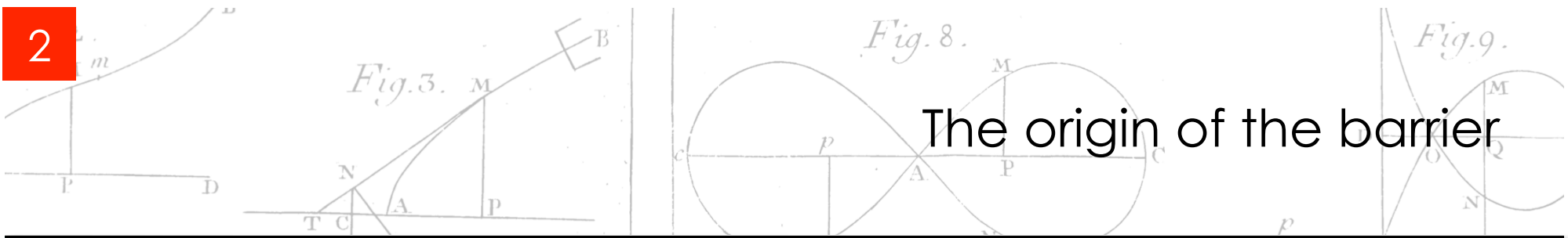


Fig. 9.

Energy landscape

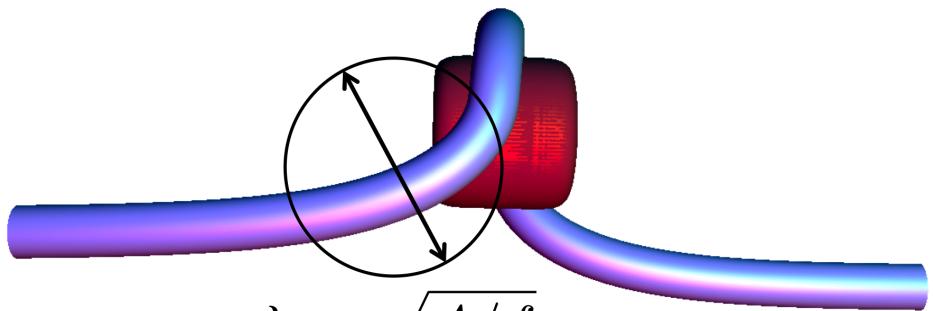
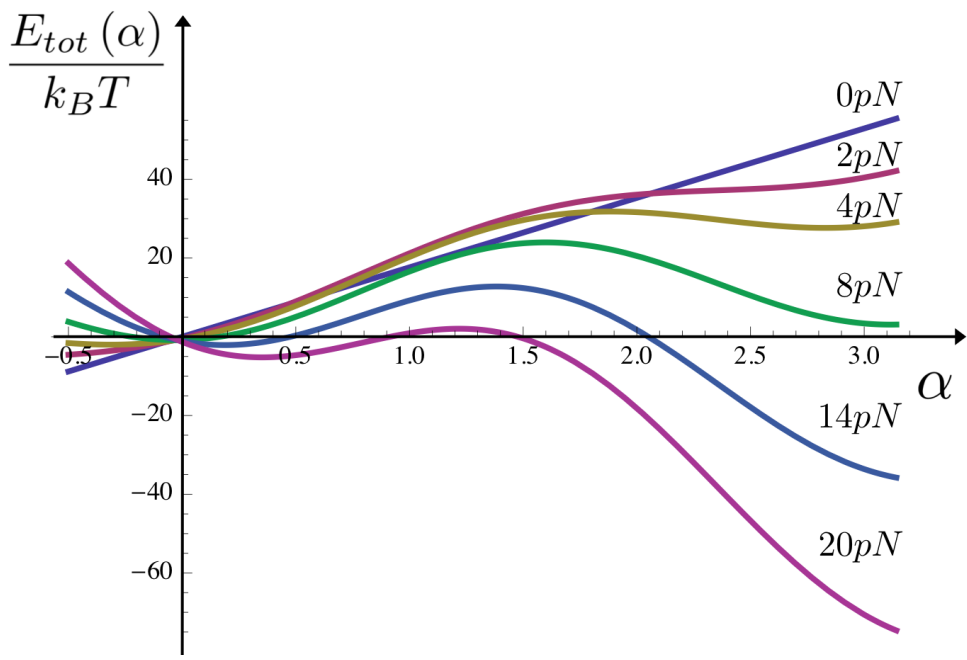
$$f = 14pN$$





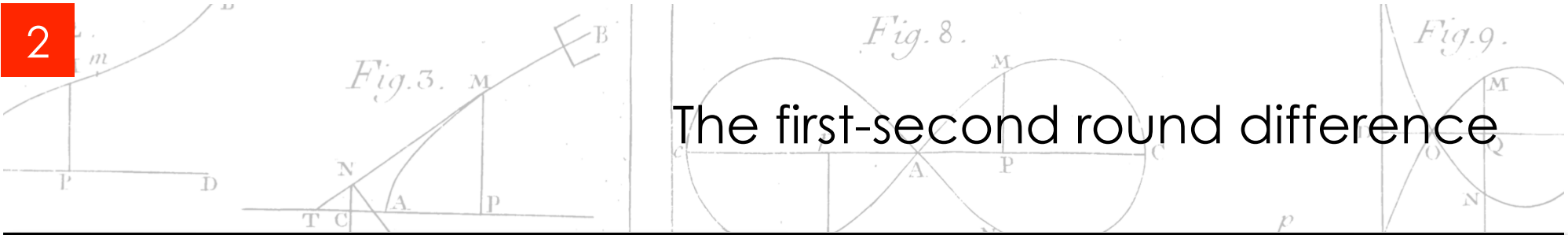
The origin of the barrier

$$E_{\text{tot}}(\alpha) \approx \underbrace{2R_0 (f_{\text{crit}} - f) \alpha}_{\text{tilting term}} - \underbrace{\frac{2}{\sqrt{3}} \sqrt{A} f \cos 2\alpha}_{\text{barrier term}}$$

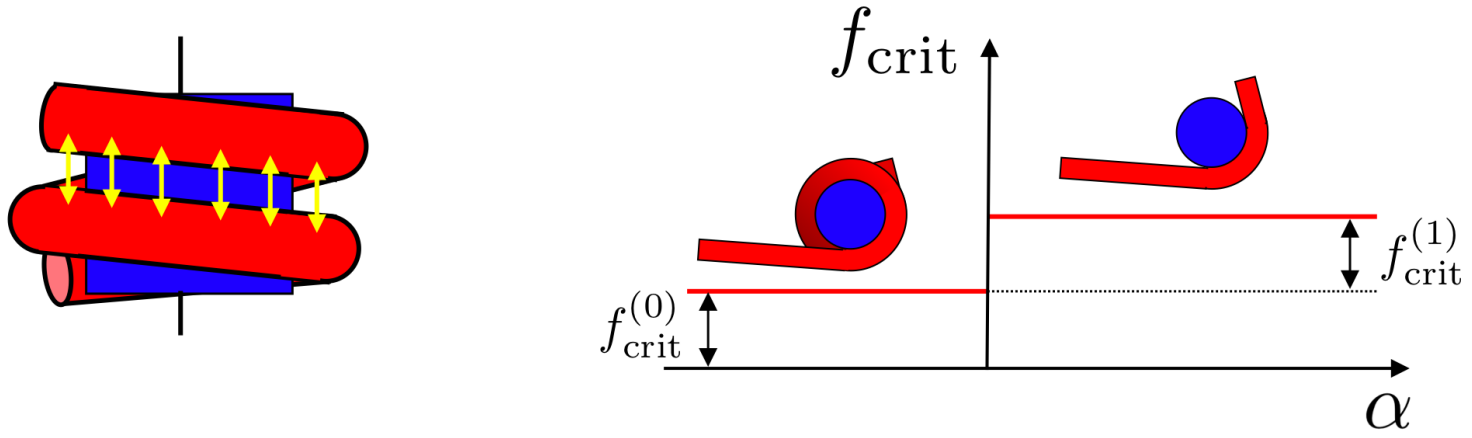


$$\lambda = \sqrt{A/f}$$

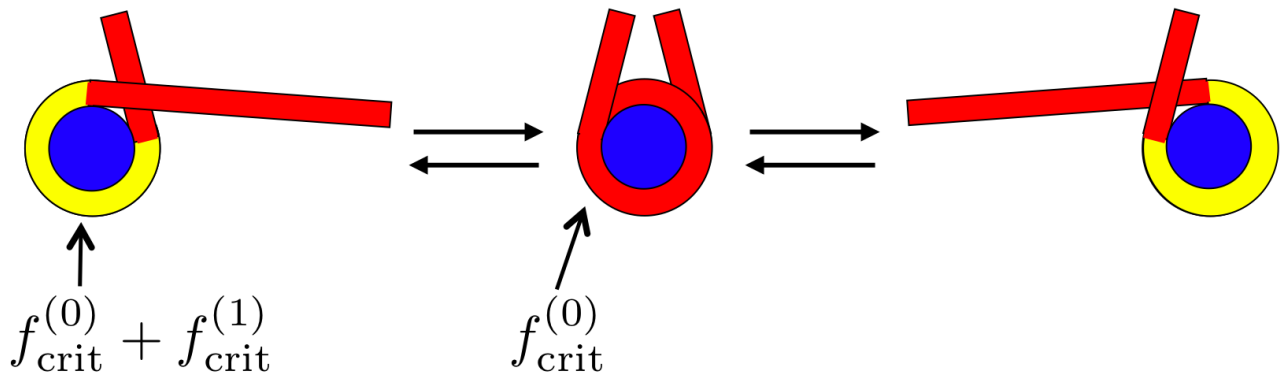
$$E_b \approx \frac{A\lambda}{\lambda^2} = \sqrt{A}f$$

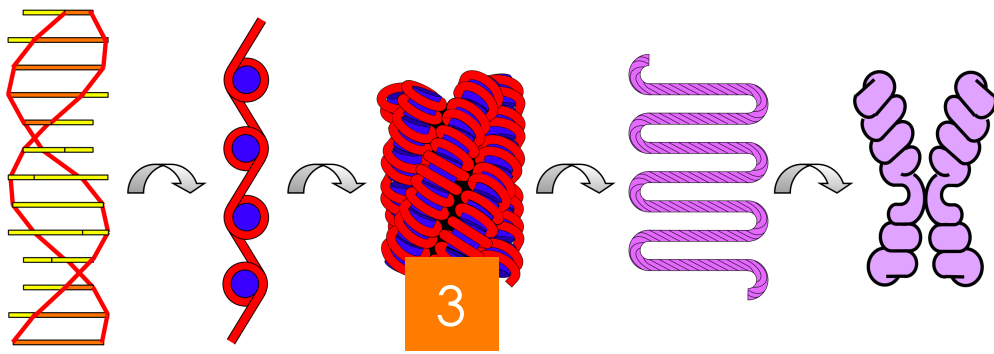


The first-second round difference

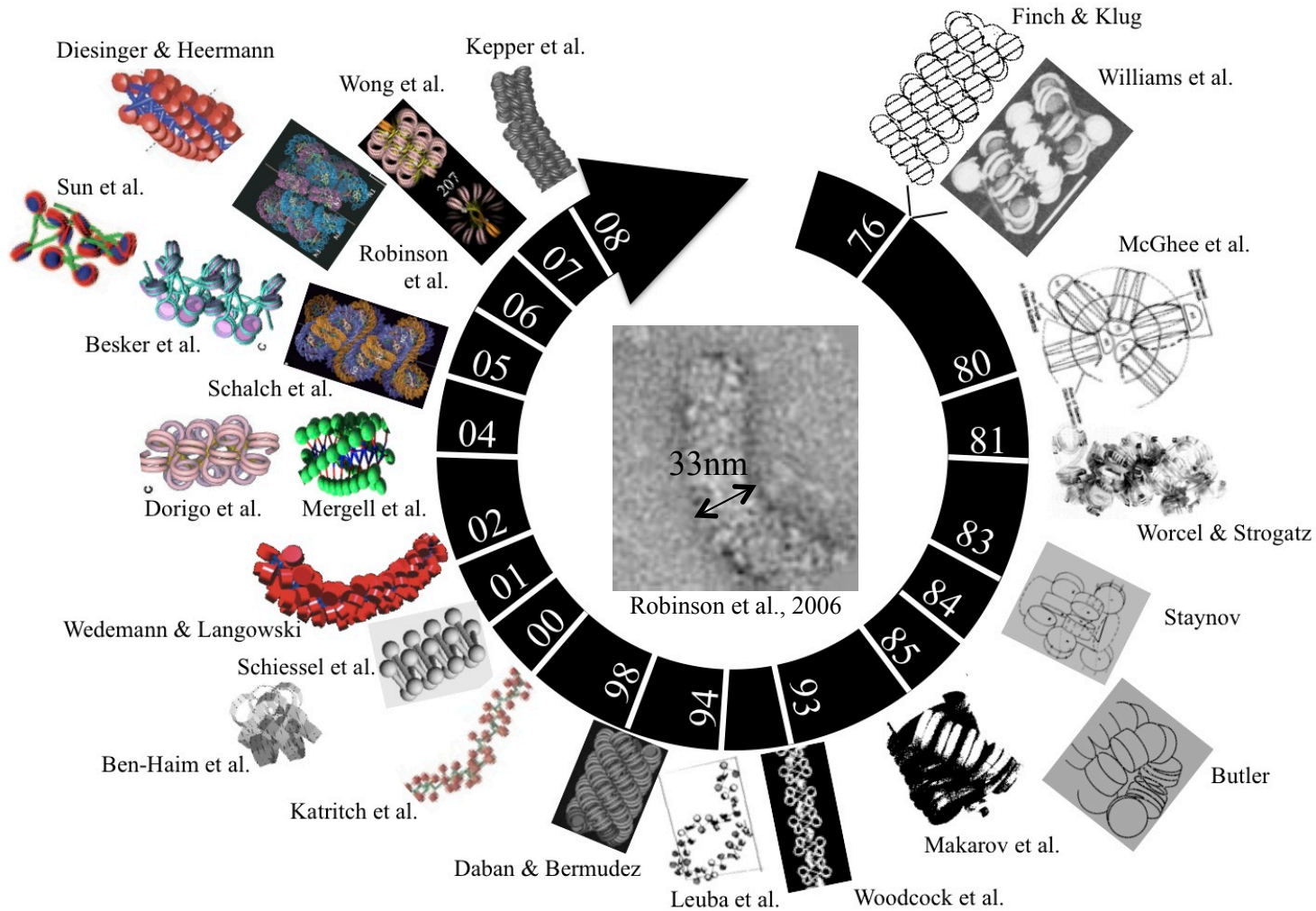


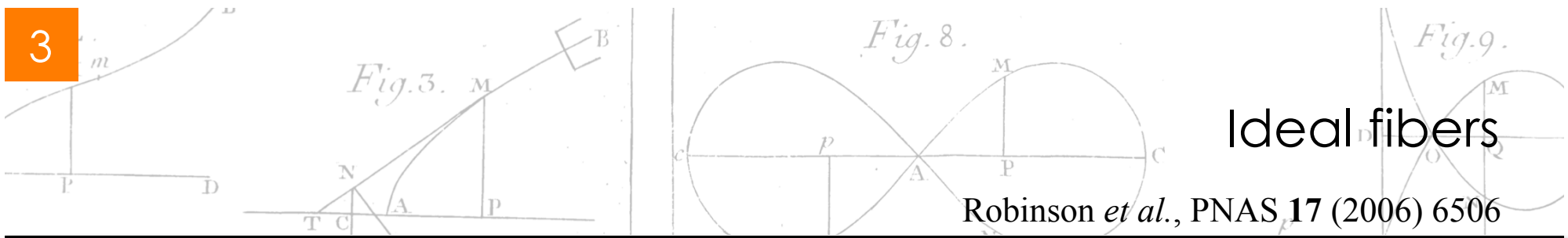
Two turn design combines accessibility and stability:





Chromatin fiber timeline

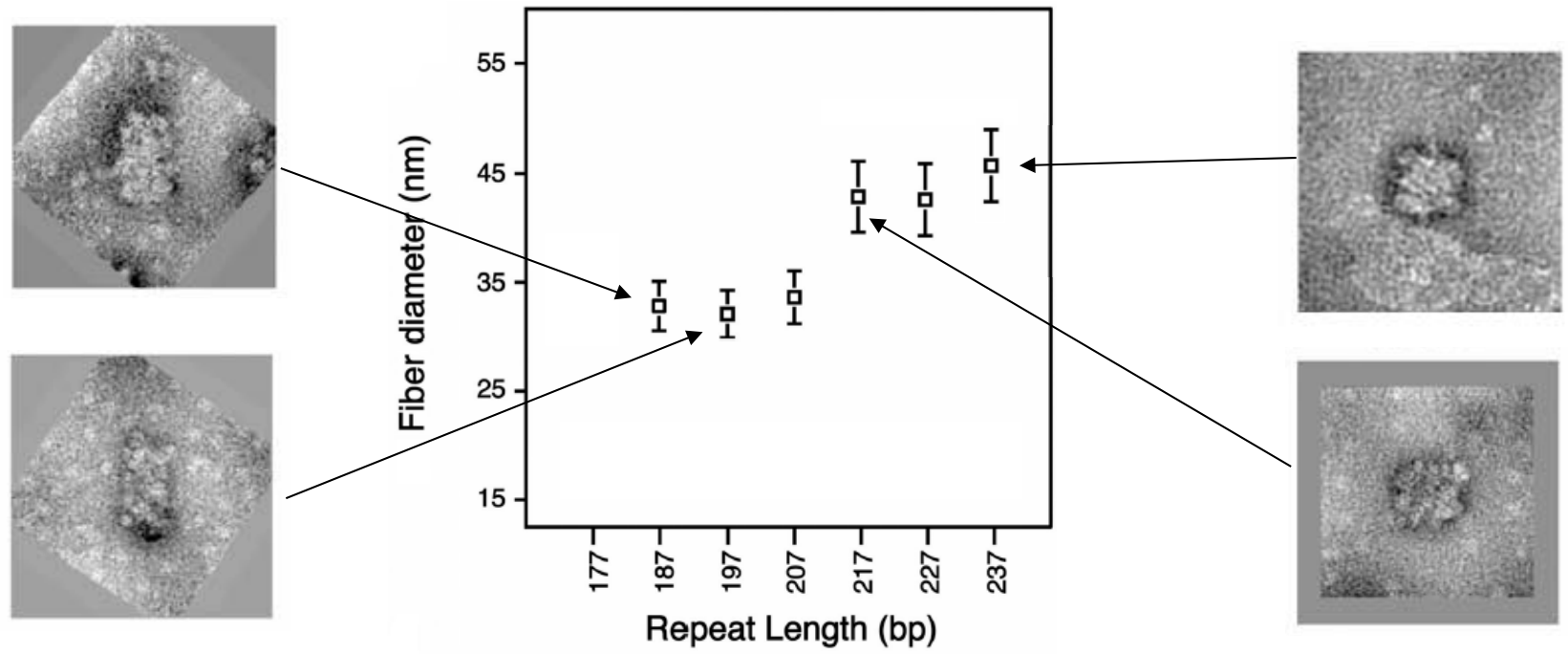
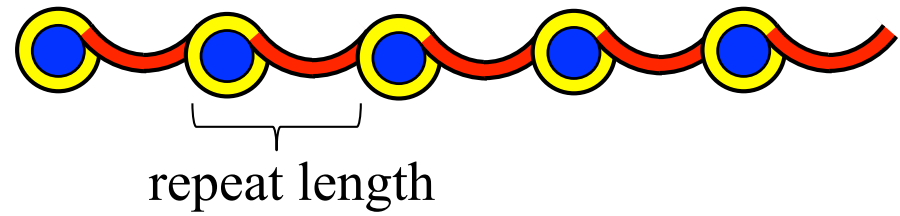


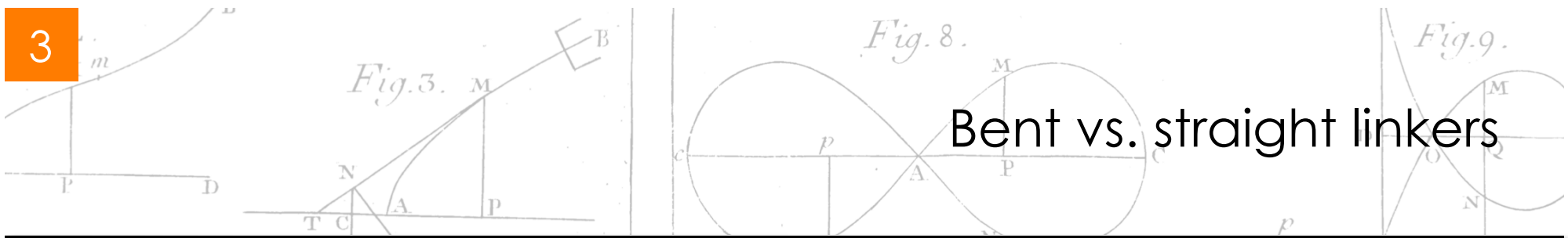


Ideal fibers

Robinson *et al.*, PNAS 17 (2006) 6506

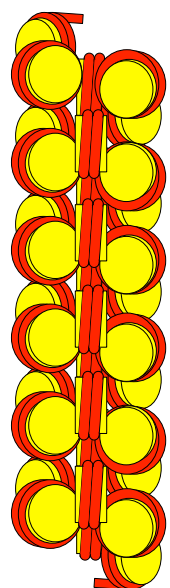
long arrays of 50 to 70 nucleosomes:





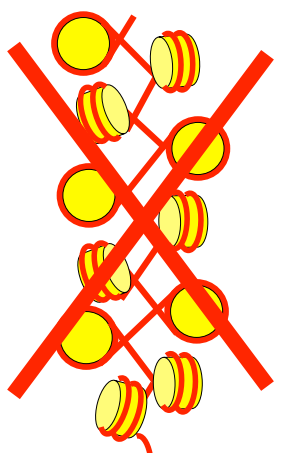
Alberts et al. "Molecular Biology of the Cell":

1994

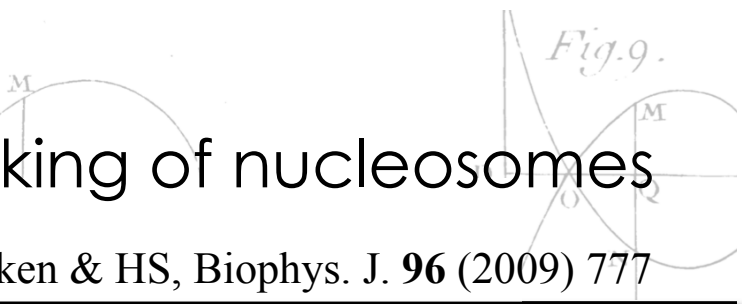
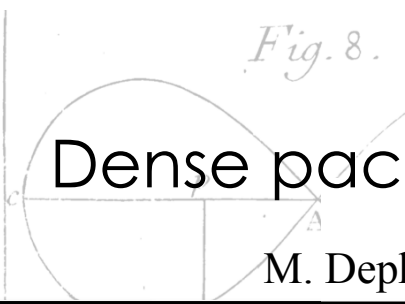
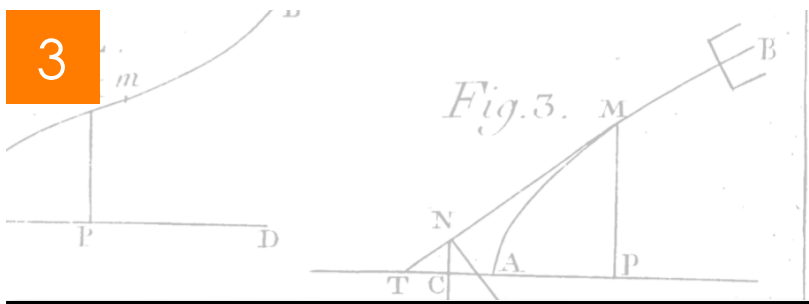


solenoid model

2002

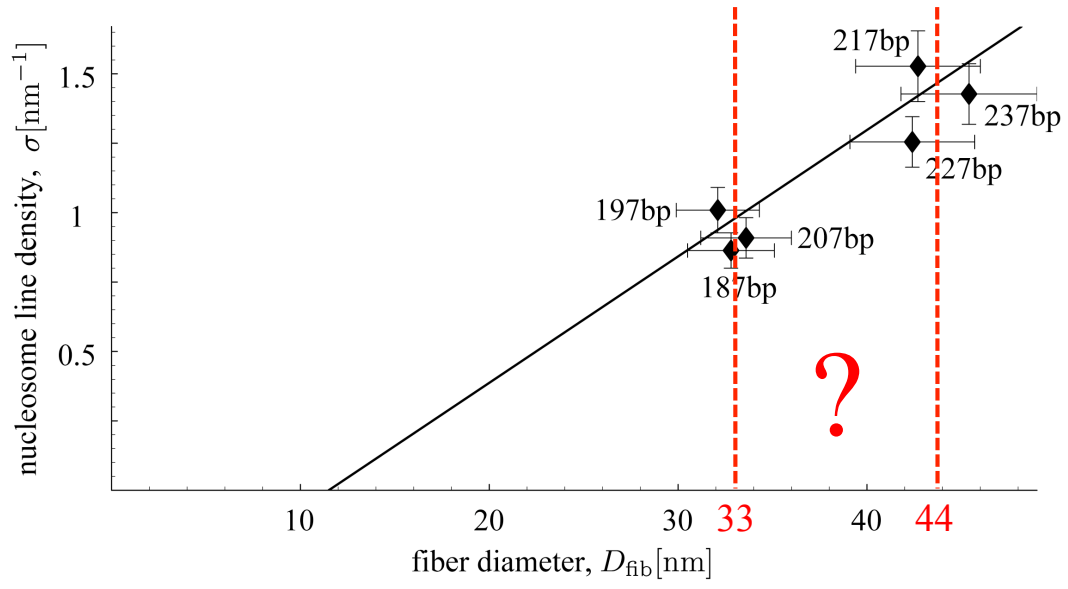
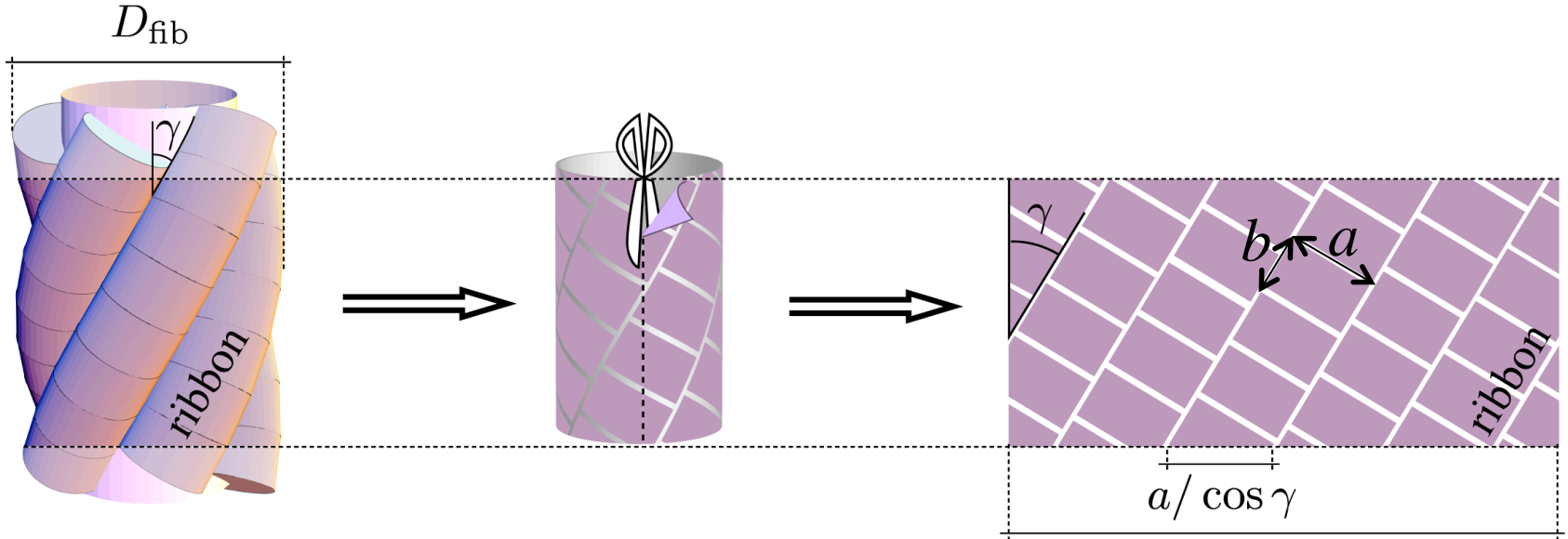


crossed-linker model



Dense packing of nucleosomes

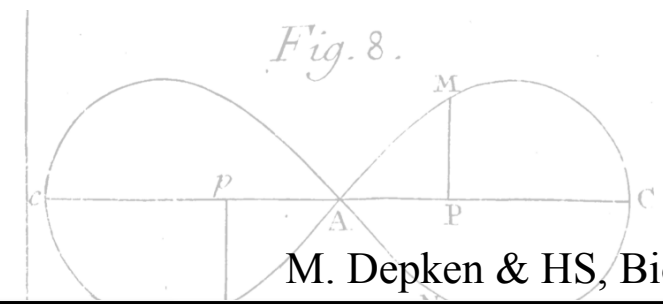
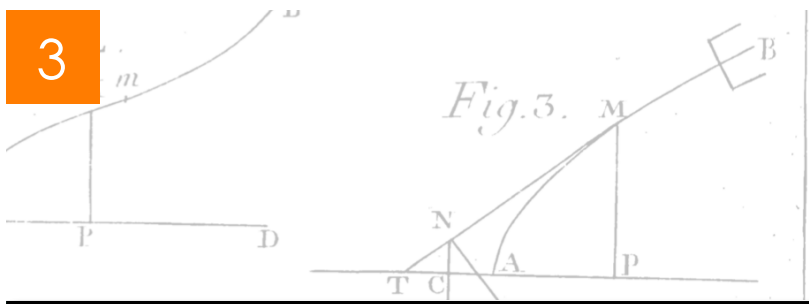
M. Depken & HS, Biophys. J. 96 (2009) 777



dense packing:

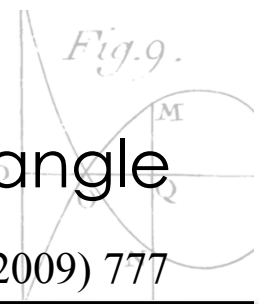
$$\sigma = \frac{\pi (D_{fib} - a)}{ab}$$

nucleosome line density

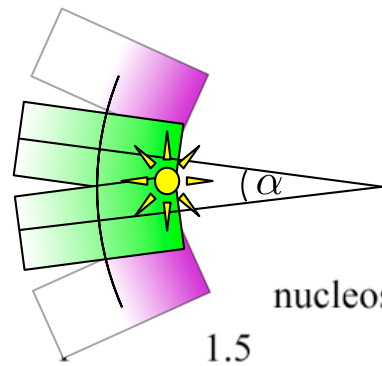
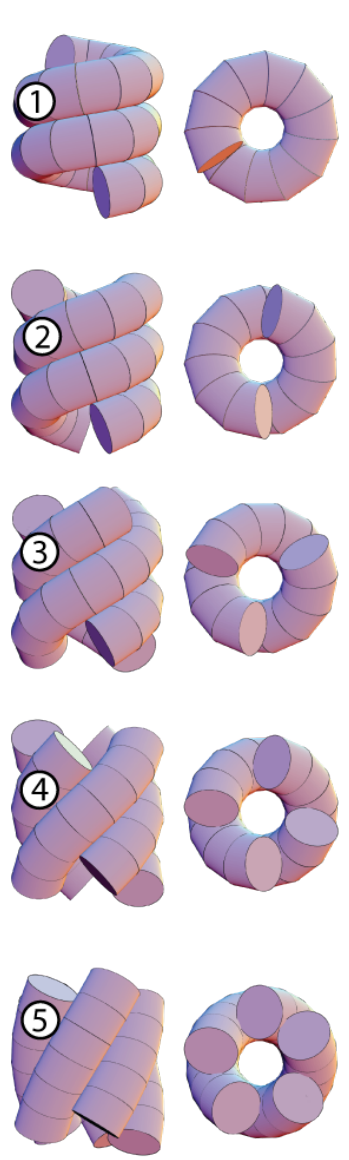


Splay angle

M. Depken & HS, Biophys. J. 96 (2009) 777

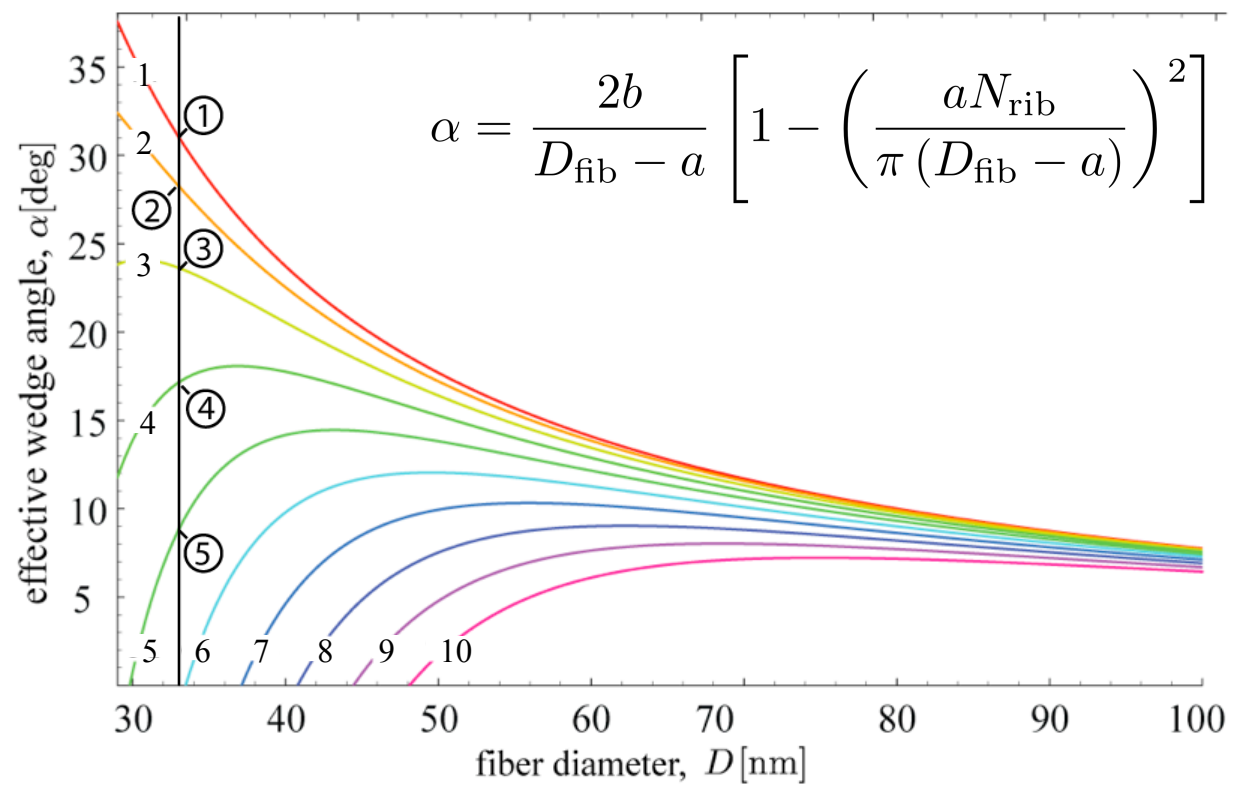


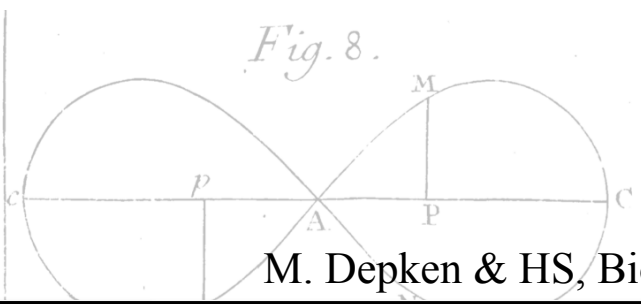
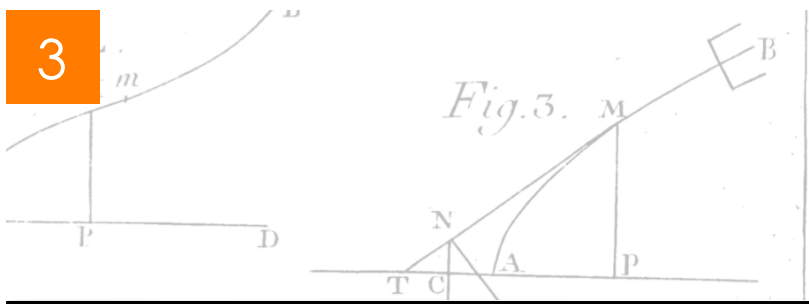
- 1976 
- 2005 
- 1985 
- 1998 
- 2007 



nucleosome line density, σ [nm⁻¹]

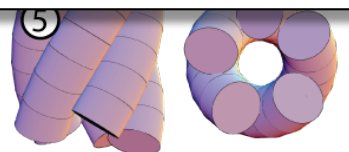
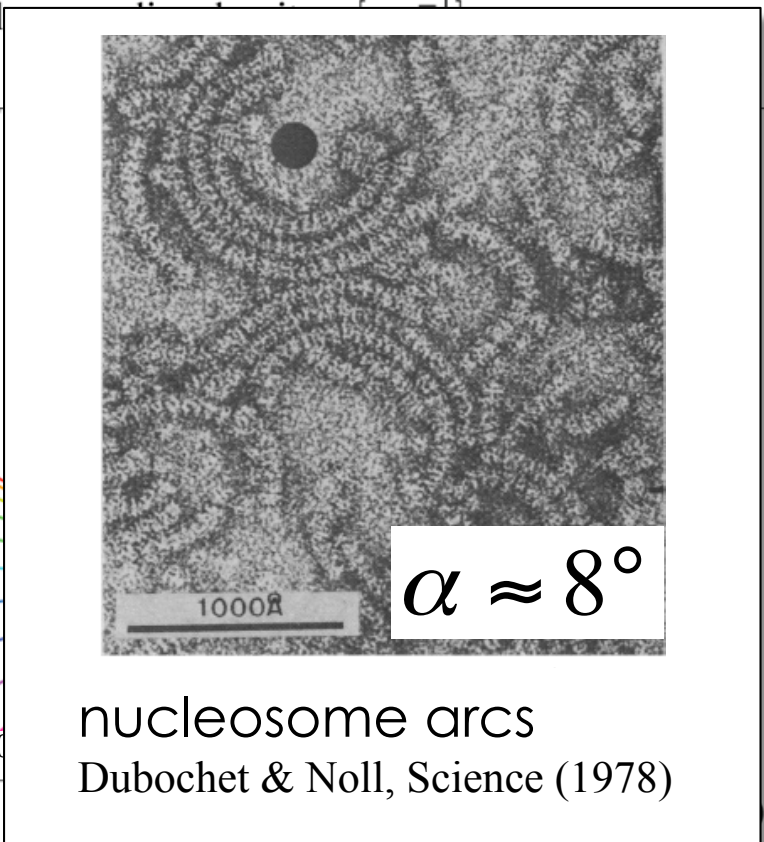
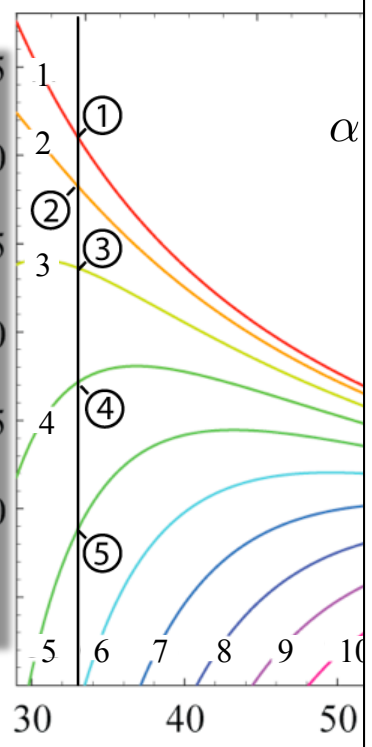
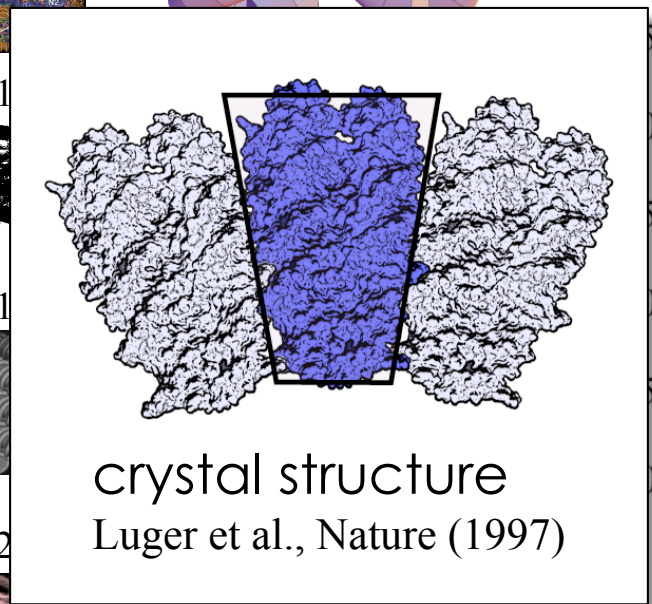
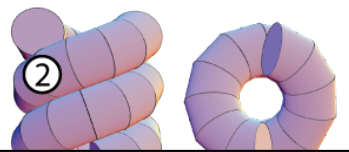
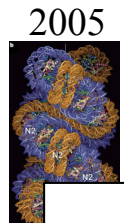
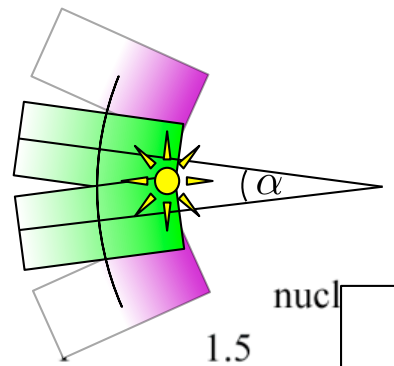
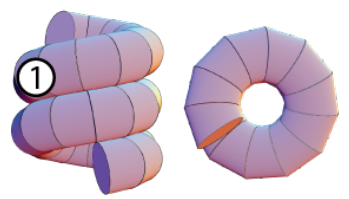
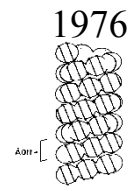
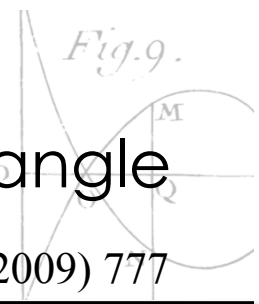
1.5 2 2.5 3 3.5 4



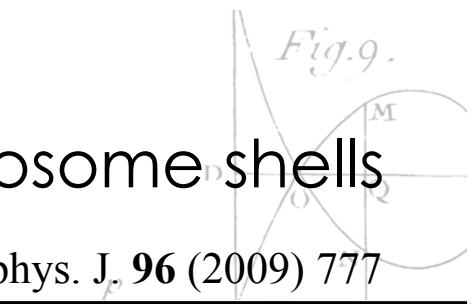
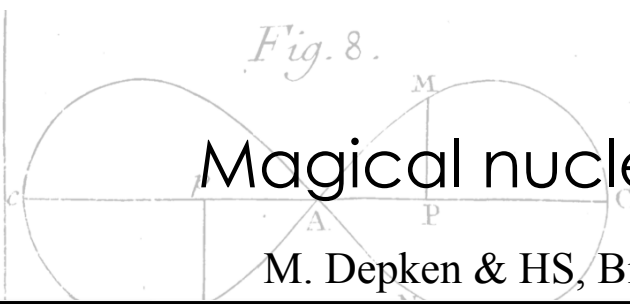
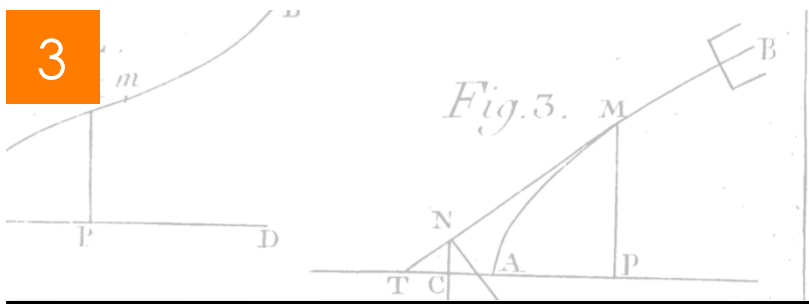


Splay angle

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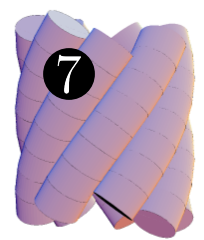
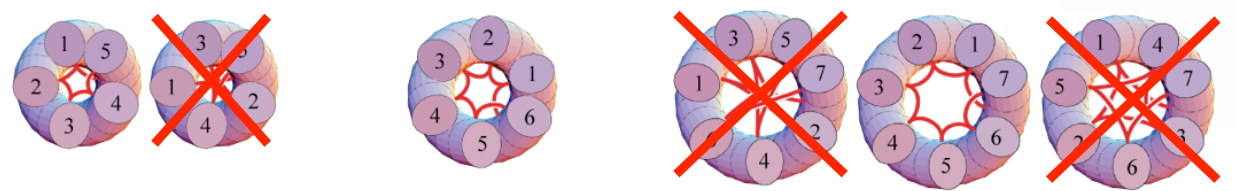
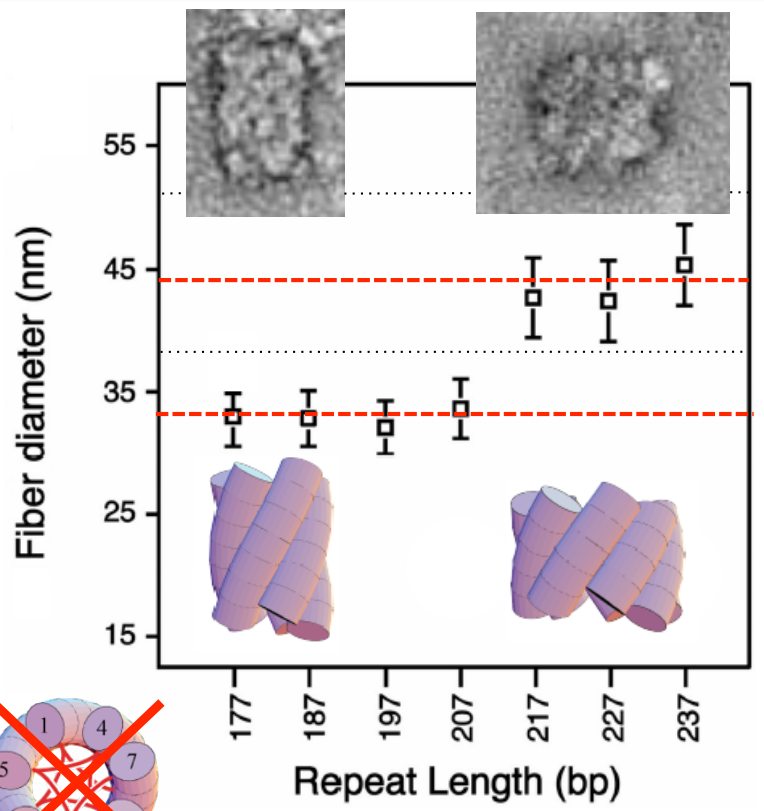
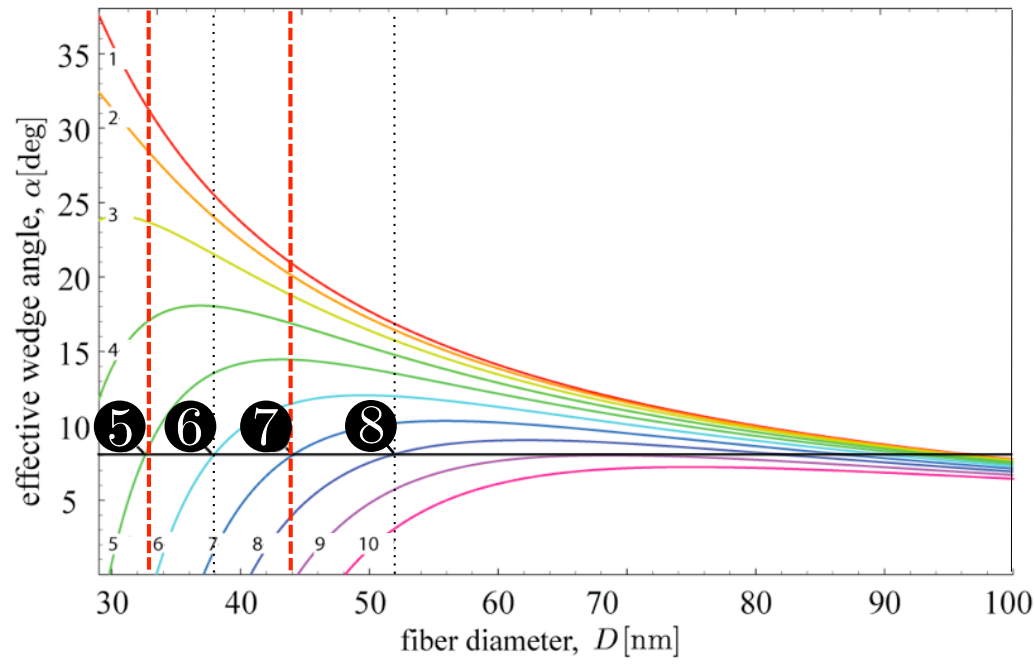


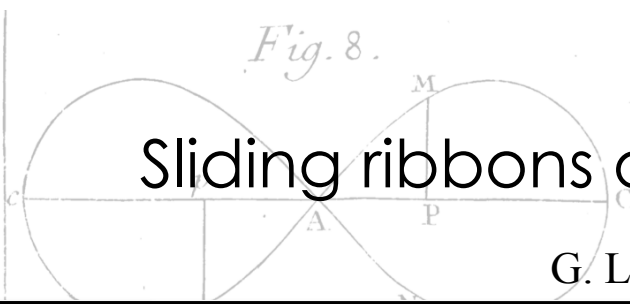
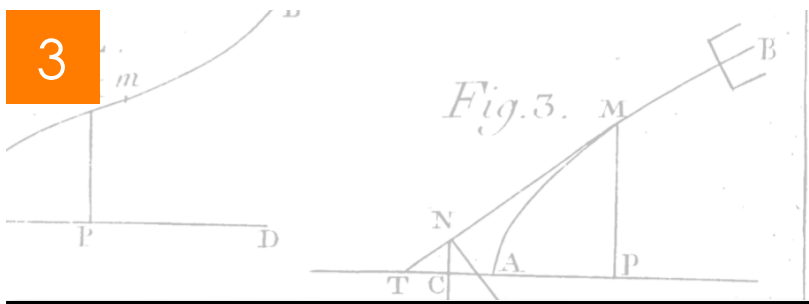
fiber diameter, D [nm]



Magical nucleosome shells

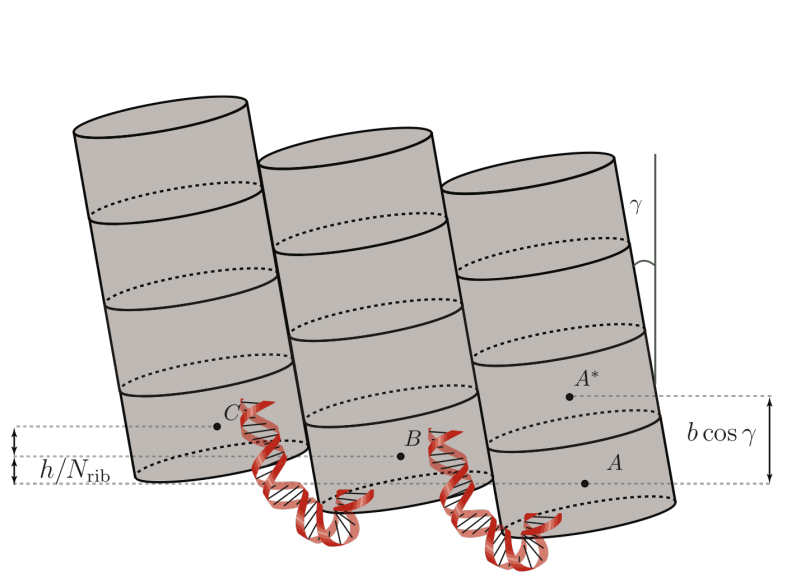
M. Depken & HS, Biophys. J. 96 (2009) 777



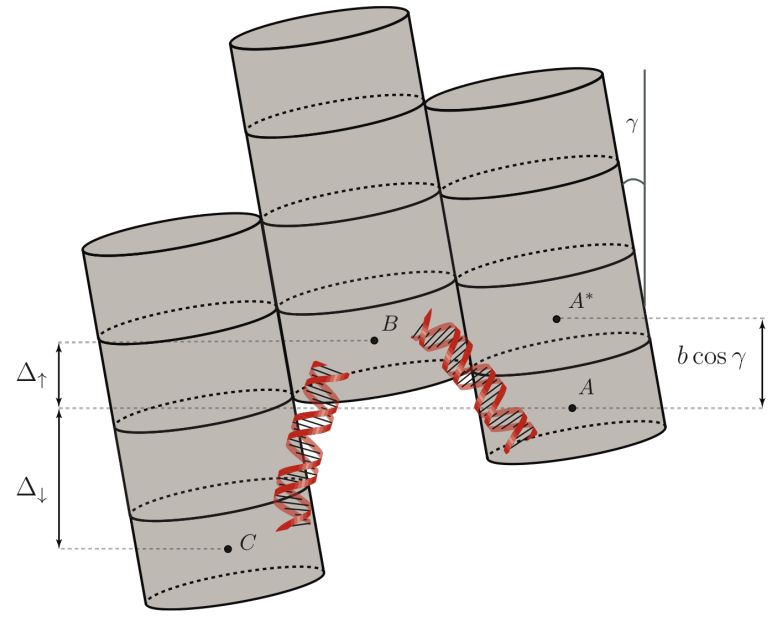
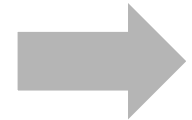


Sliding ribbons out-of-register

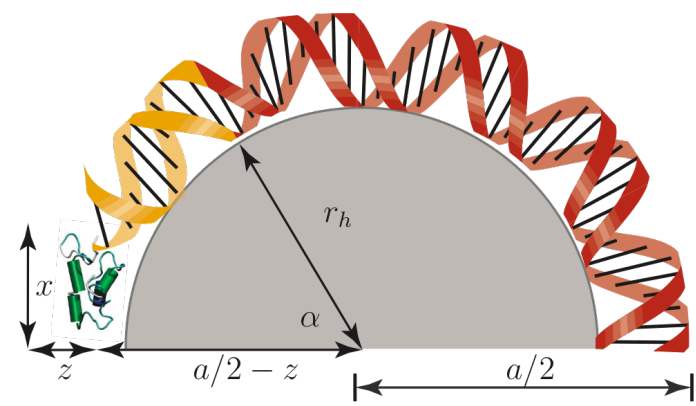
G. Lanzani & HS, submitted



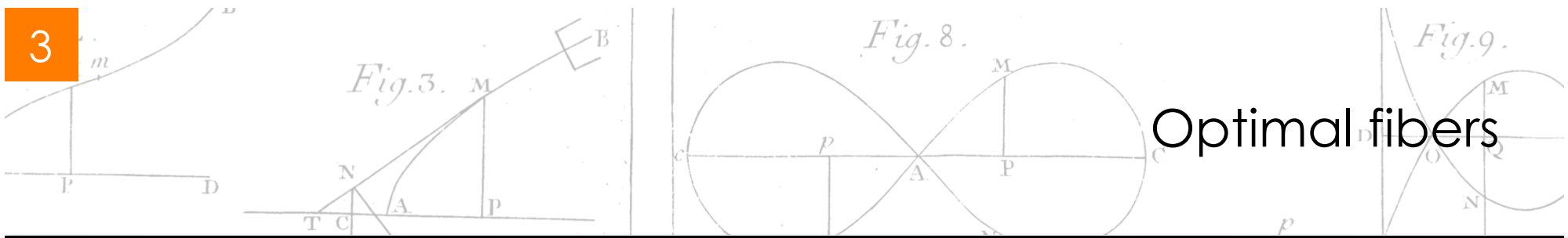
$$\approx 40k_B T$$



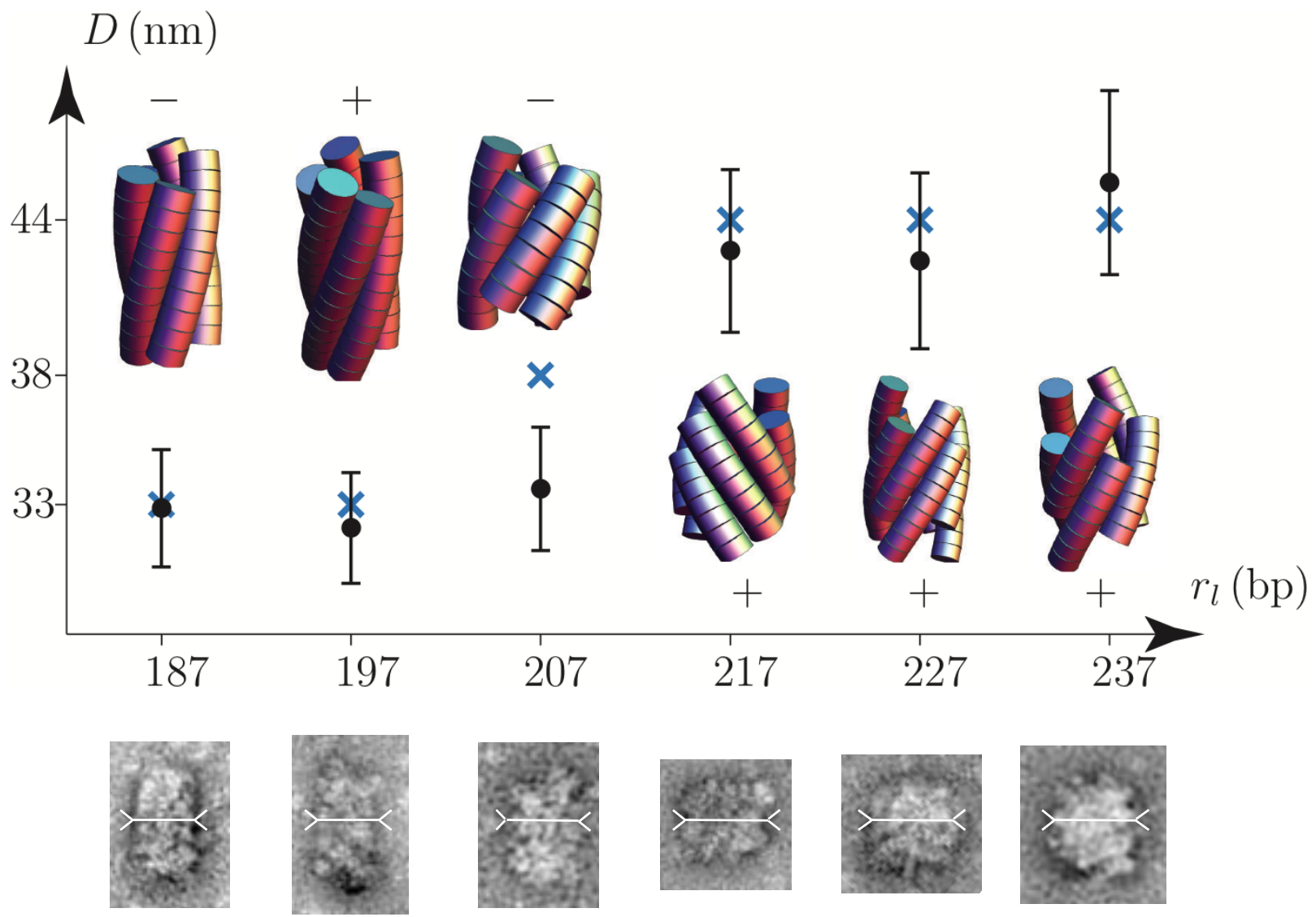
$$\approx 1k_B T$$



3



Optimal fibers



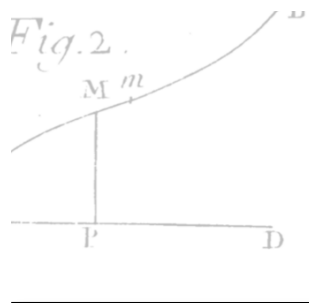


Fig. 3.

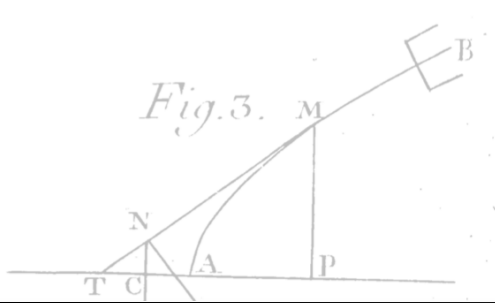


Fig. 8.

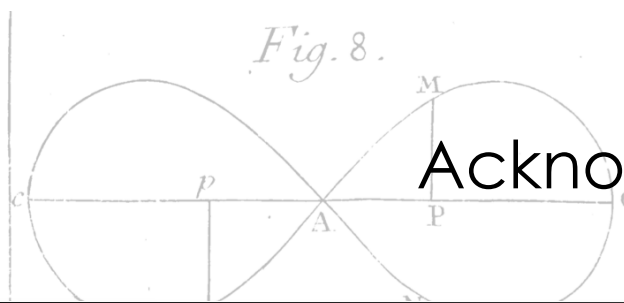
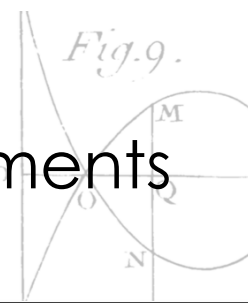
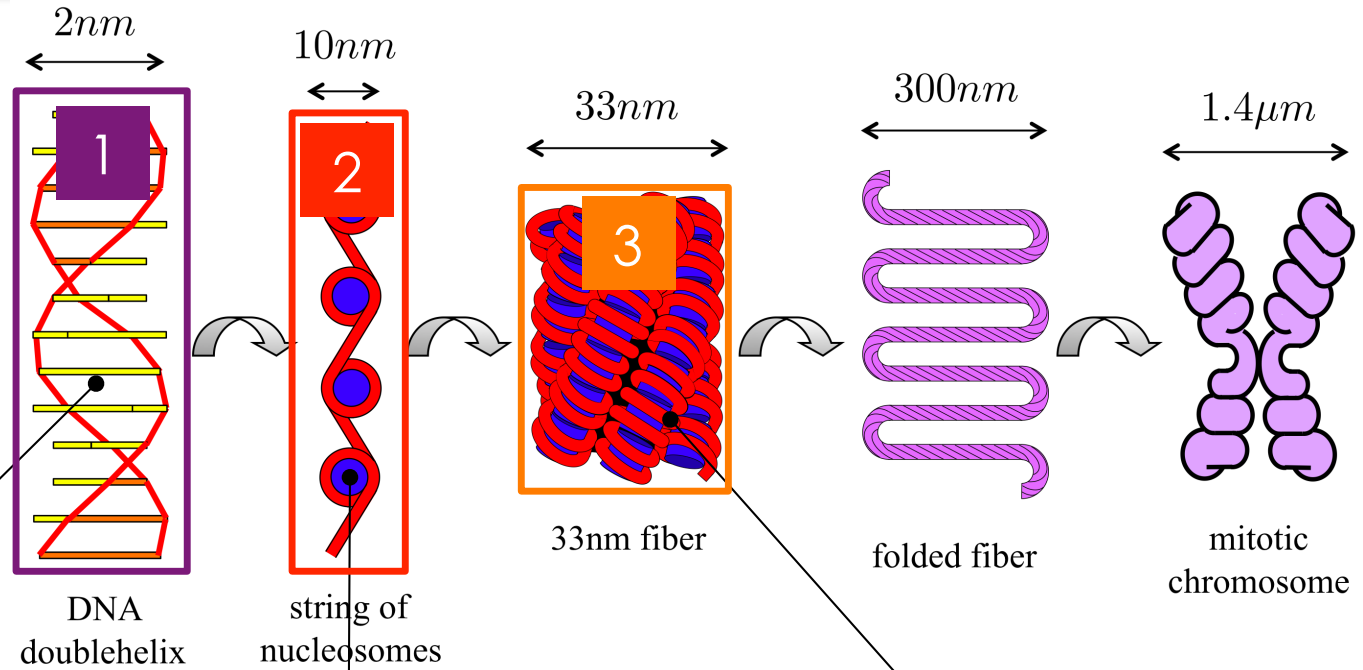


Fig. 9.



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VU Amsterdam