

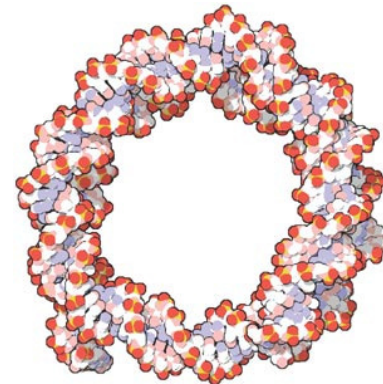
High flexibility of DNA on short length scales probed by atomic force microscopy

**Wiggins P. A. et al.
Nature Nanotechnology (2006)**

**presented by Anja Schwäger
23.01.2008**

Outline

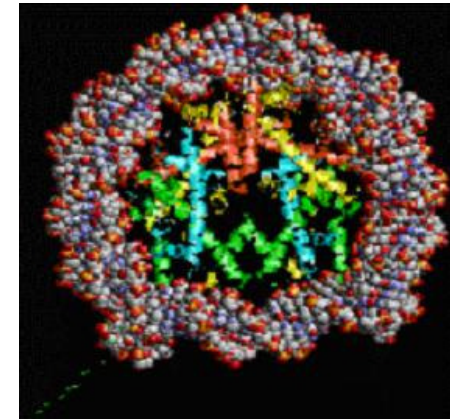
- Theory/Background
 - Elasticity of DNA
 - Aim of study
- Methods
 - AFM
 - Data analysis
- Results
 - New model for DNA elasticity
 - Comparison with other results
- Outlook
 - Improvement of model
 - Summary



DNA mechanics

DNA elasticity

- Bending and twisting properties of DNA
 - ⇒ determine DNA conformation and dynamics
- Elastic properties of DNA
 - double helix relatively stiff for bending
 - unlikely to form sharp bends, high energy needed
- Biological importance of sharp DNA bends on short length scales
 - chromatin
 - packaging of DNA in capsid
 - protein interactions (gene regulation)



Nucleosome

<http://www.jyi.org/research/re.php?id=419>

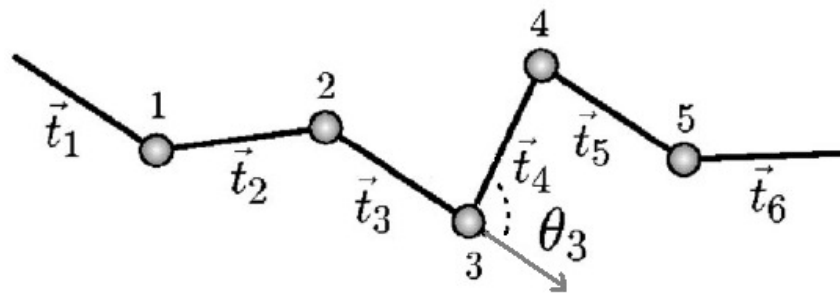


Lac Operator

Swigon, PNAS, 2006

Worm-Like Chain (WLC)

- **Elasticity model** for the description of **semiflexible polymers**
- Molecule as a thin rod with constant and isotrop elastic behaviour
- Approximation (discrete segments):
 - Conformation described as a chain of segments
 - Each segment with classical elastic energy cost for bending (Hooke's law)
- WLC is this approximation in the limit of zero segment length



modified according to Wiggins and Nelson, Phys Rev E, 2005

Worm-Like Chain (WLC)

- Important parameter of the WLC:

Persistence length of a polymer

Measure for molecule stiffness / the resistance of the chain to bending

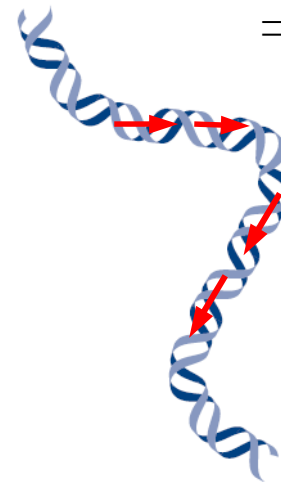
(length over which correlation of tangent vector directions is lost
 \approx distance that separates bends)

short persistence length
 \Rightarrow more flexible



long persistence length

\Rightarrow stiffer



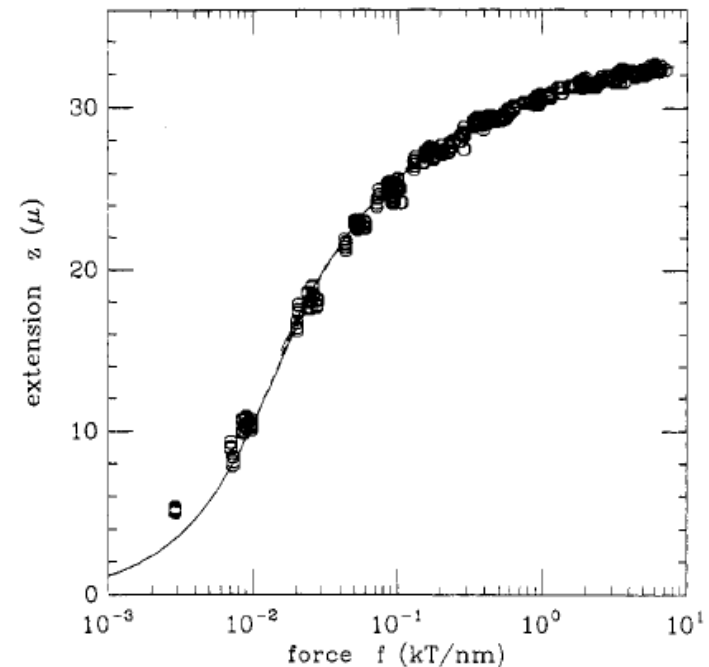
Podgornik, Nature Nanotechnology, 2006

WLC description of DNA

Example: Stretching of DNA



- Simple description of the behaviour of a polymer chain as response to extension force
 - reduction of entropy with straightening
- Length scale longer than persistence length
 - results for $L = 32.8 \mu\text{m}$
 - persistence length of double helix DNA $L_p \approx 50 \text{ nm (150 bp)}$



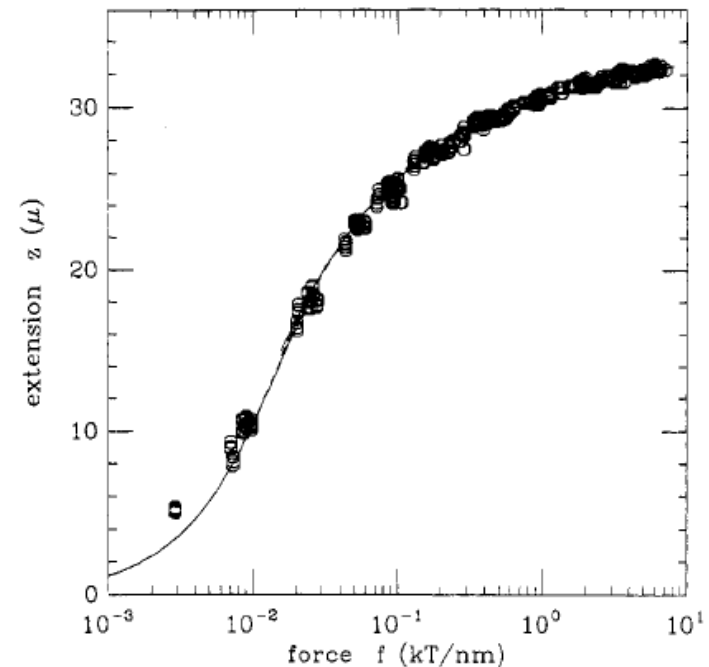
Marko and Siggia, Macromolecules, 1995

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Marko and Siggia, Macromolecules, 1995

WLC description of DNA

Bending energy of a conformation

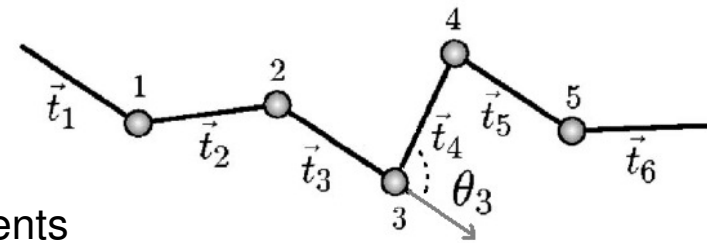
Chain of segments of length ℓ with **harmonic energy** cost per segment:

$$E_{\text{WLC}}(\theta) = \frac{1}{2} k_B T (L_P/\ell) (\theta_i)^2$$

θ : bending angle between segments

L_P : persistence length of DNA

$k_B T$: thermal energy



modified according to Wiggins and Nelson, Phys Rev E, 2005

Energy rises quadratically with bending angle

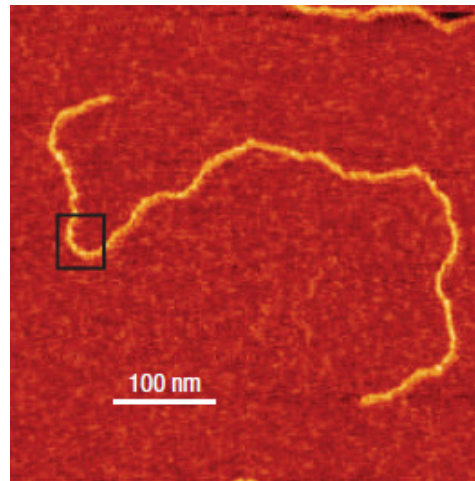
⇒ **Boltzmann distribution**: large bending angles will occur rarely

Motivation of experiments

- Previous studies of DNA mechanics focus on longer length scales
 - Cyclization studies of **short DNA fragments**: short circles form much more easily than expected from WLC theory
 - **Test WLC** model on short length scales (few nanometers)
- ⇒ Measure model predicted **bending angle frequencies** directly in equilibrium DNA conformations
- ⇒ Frequencies of bending angles correspond to **bending energy function**

Experimental approach

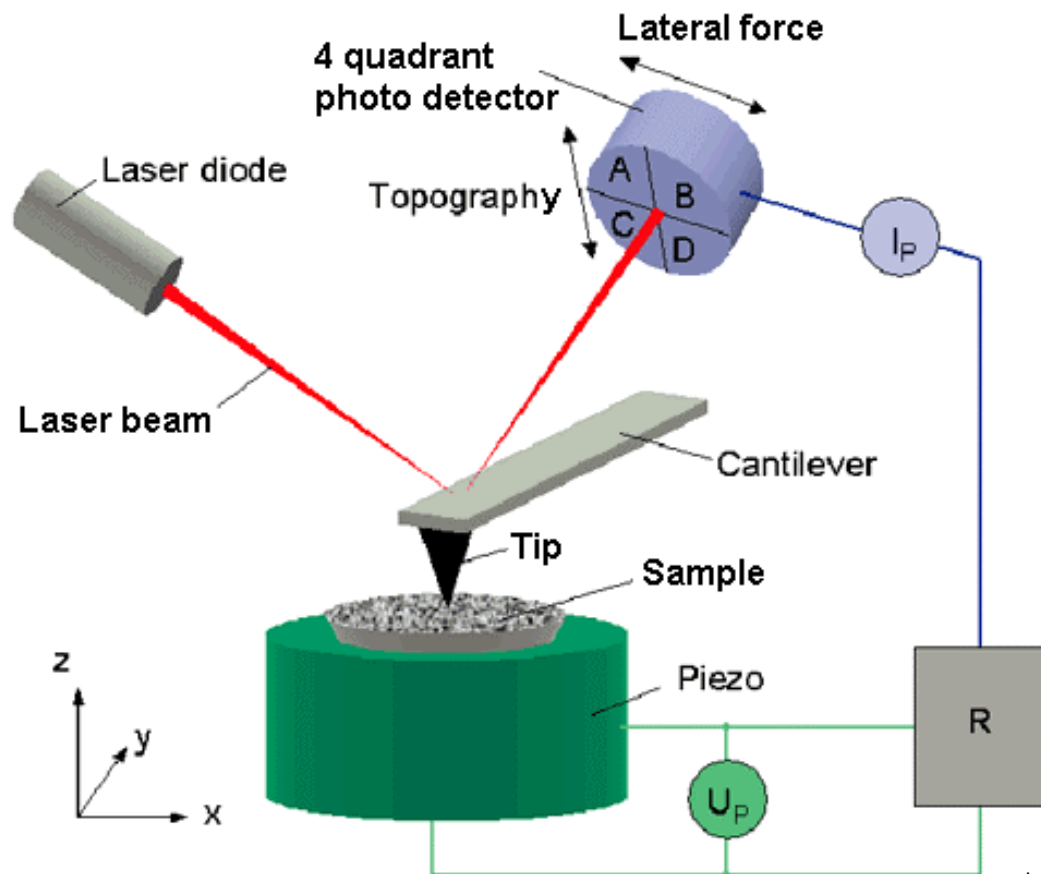
- High-resolution **atomic force microscopy**
- 2743 bp **linear double-stranded DNA** with random sequence, **adsorbed to surface**
- **Two-dimensional** movement possible
- Assume **equilibrium conformation** of adsorbed molecules



Wiggins, Nature Nanotechnology, 2006

Atomic Force Microscopy (AFM)

Principle



Characteristics

- topographical images
- near angstrom resolution (0.1 nm)

Biological application

- imaging of living cells
- measurement in physiological buffers

<http://www.nanomat.de/datenblaetter/42000.htm>

Theory

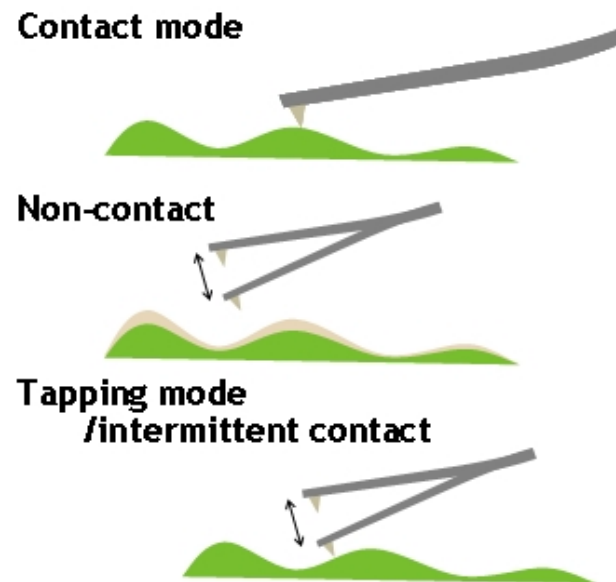
Methods

Results

Outlook/Summary

AFM operation modes

Typical modes



<http://en.wikibooks.org/wiki/Nanotechnology/AFM>

- Tapping mode usually used for soft biological materials

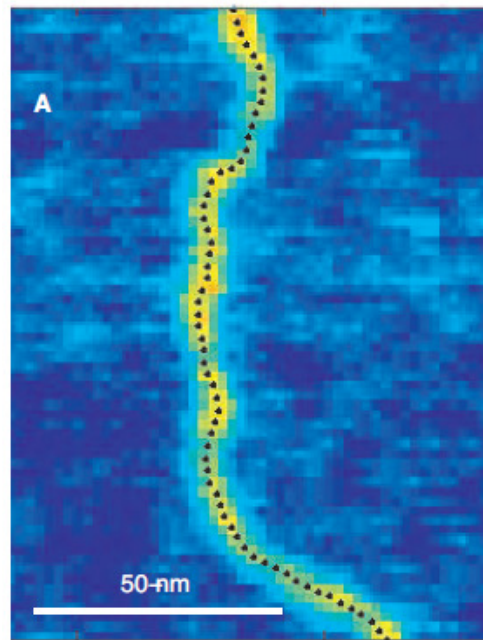
Experimental and simulated AFM data

- Linear DNA molecules adsorbed to mica (silicate surface) with different concentrations of Mg^{2+} ions
 - AFM measurement in tapping mode **in air**
 - height profile of equilibrium conformations
 - Simulation of AFM data on the basis of WLC
 - generation of WLC conformations, 2D chains with bends
 - bending angle distribution according to WLC prediction
 - conversion to simulated AFM images
 - same analysis as experimental conformations
- ⇒ Exclude that conformations are artifacts of AFM measurement

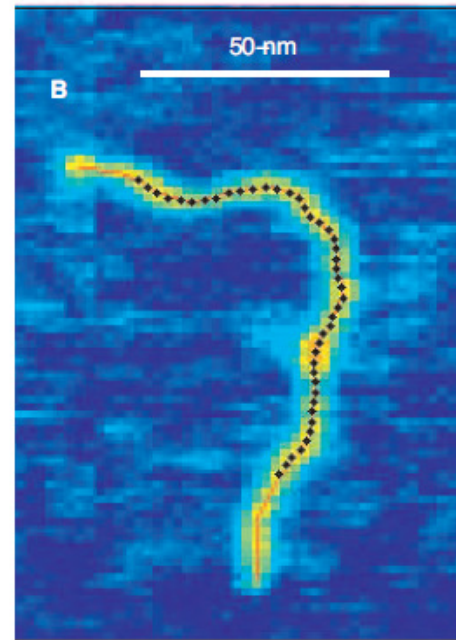
Image analysis

Automatic tracing of DNA contours

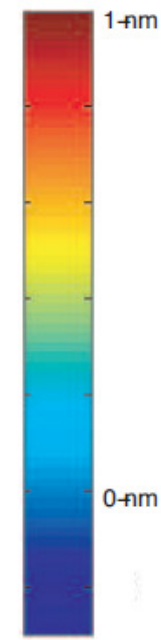
- Set points with a distance of 2.5 nm on the contour
- Length ℓ of segments 2.5 nm



experimental image



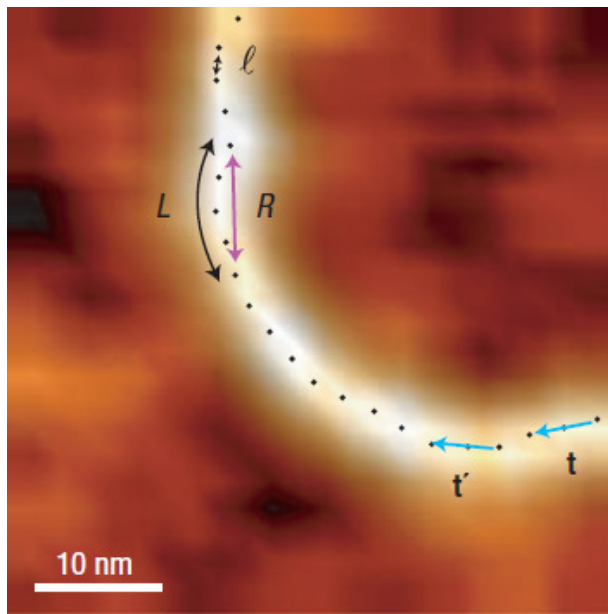
simulated image



Wiggins, Nature Nanotechnology, 2006

Measurement of angles

- Two points of chain separated by the **contour length L**
- **Tangents** of these points t and t'
- Angle between two tangents
- Measured for all points on complete chain and different L



Example: $L = 3 \times 2.5\text{ nm}$
contour length between t and t'

Wiggins, Nature Nanotechnology, 2006

Statistical evaluation

- **Angle Distribution $G(\theta; L)$:**
Probability distribution of angles between tangents
 - angles between tangents t_i and t_{i+1} separated by L measured for many molecules
 - histogram of angles
- Curve measures the **bending energy $E(\theta)/k_B T$** ($E(\theta)$ in units of $k_B T$) of the segments of length L

Bending angle distribution:

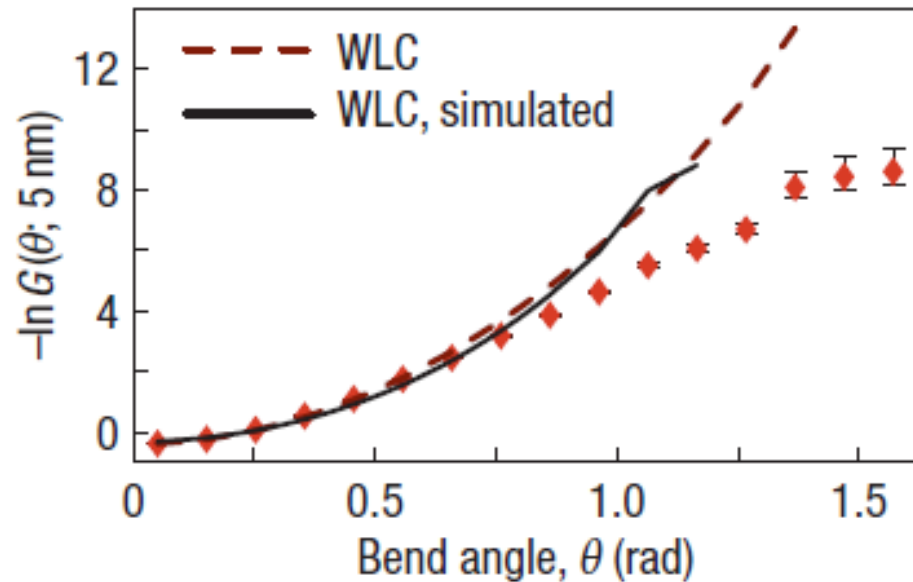
$$g(t_{i+1}|t_i) = q^{-1} \exp[-E(\theta_i)/k_B T]$$

Negative logarithm:

$$-\ln(g) = \ln(q) + E(\theta_i)/k_B T$$

Results

Dots: measured angle distribution for $L = 5$ nm (negative logarithm)



Wiggins, Nature Nanotechnology, 2006

- Distribution calculated from theoretical model (WLC) does not match the measured data for large angles
- Values of measured data are much lower (= higher frequency of large angles)
- Simulated WLC data match theoretical curve

Results

- Segments with large bends much more frequent than predicted by WLC
 - large bends (≥ 1.1 rad) 30 times more frequent
 - Elastic energy for high bending angles must be lower than predicted
 - Energy function is almost linear
- ⇒ Hooke's law does not apply for the elastic energy on small length scale

Other experiments

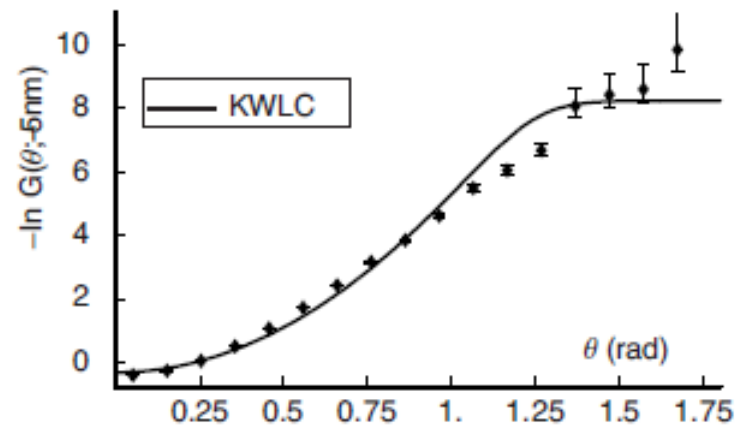
- Cyclization studies by Cloutier and Widom
- Sharply looped DNA highly bendable and also **highly twistable**, DNA more flexible than predicted by models
- Theories of DNA flexibility:

Melted bubbles, single-stranded regions with high flexibility, facilitate loop formation

Kinkable worm-like chain

Extension of the WLC that includes spontaneous kink formation when curvature is high

⇒ Local alternative conformations with high flexibility



Wiggins, Nature Nanotechnology, 2006

Development of new model

- Equation that fits measured data
- **Linear ‘sub-elastic chain’ (LSEC) model**

$$E_{\text{LSEC}}(\theta) = \alpha |\theta| k_B T$$

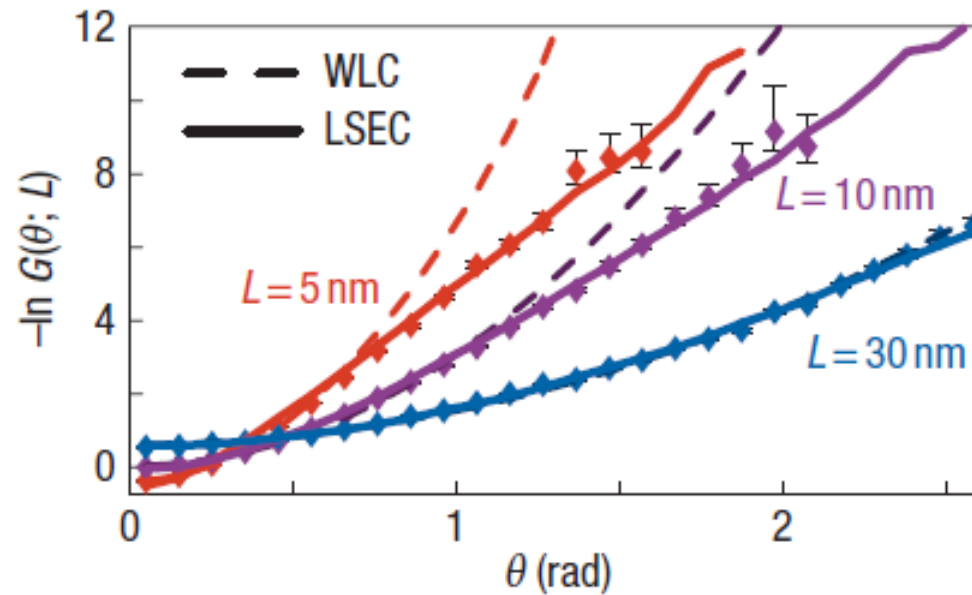
θ : bending angle between segments

α : dimensionless constant, dependent on ℓ

$k_B T$: thermal energy

Non-harmonic (linear) energy function, “softer“ for sharp bending

LSEC on long scale

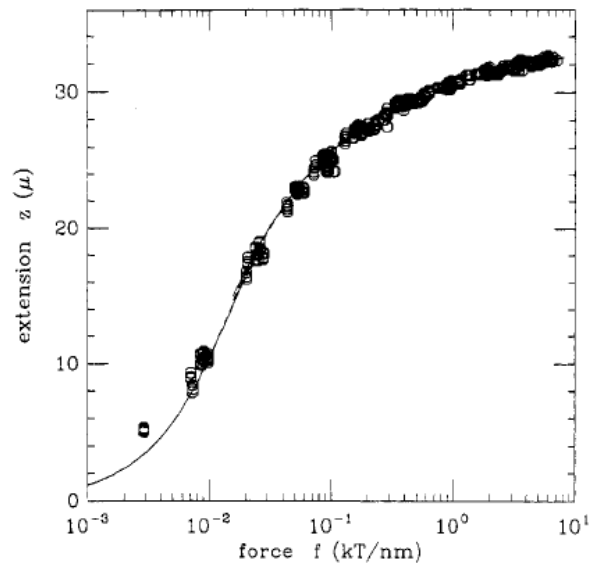


Wiggins, Nature Nanotechnology, 2006

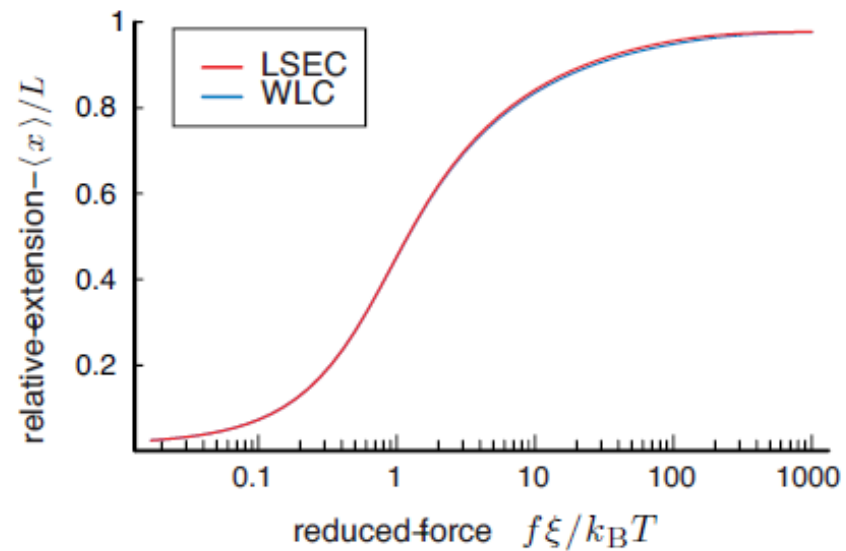
- WLC describes behaviour for **long** contour length correctly
- LSEC model fits data on **all** length scales

LSEC on long scale

- Application of LSEC on other experimental data
- Force-extension experiment correctly described by LSEC



Marko and Siggia, Macromolecules, 1995



Wiggins, Nature Nanotechnology, 2006

Outlook

Improvement of new model

- Include twisting properties in LSEC model
- Test sequence dependence of DNA elasticity

Application

- Relevance of model to other semiflexible biopolymers
 - Is DNA the only polymer with higher elasticity on short scales?

Criticism

- Further/other experiments that support findings
(Exclude that results are an artifact of the measurement)

Summary

- DNA flexibility on short length scales is important in many biological processes (e.g. nucleosome formation).
- Using high-resolution AFM the bending behaviour of DNA in such scales can be measured directly.
- Sharp bends in DNA on short length scales are much more frequent than predicted by the WLC model.
- A simple model can describe both the bending behaviour of DNA on short scales and the long-length scale observations (force-extension curve of DNA)

Literature

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Additional figures:

- <http://www.jyi.org/research/re.php?id=419>
- <http://www.nanomat.de/datenblaetter/42000.htm>
- <http://en.wikibooks.org/wiki/Nanotechnology/AFM>

**...Thank you for
your attention!**