

Single-Molecule Manipulation Experiments of Biological Molecules III: Applications to Biological Systems

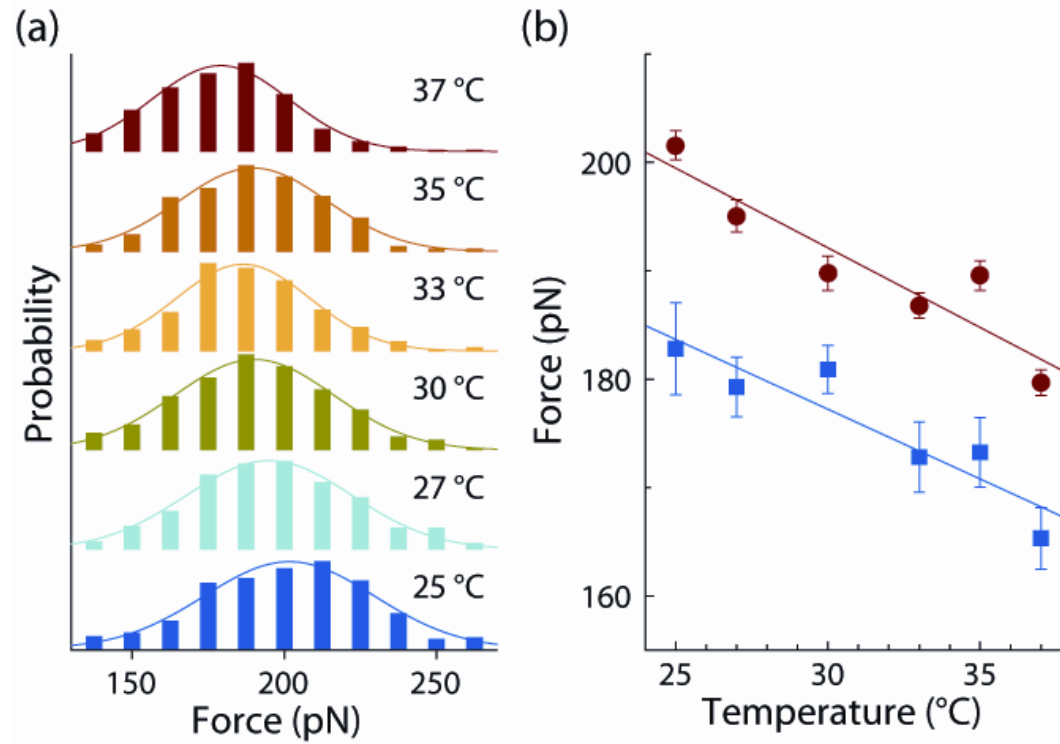
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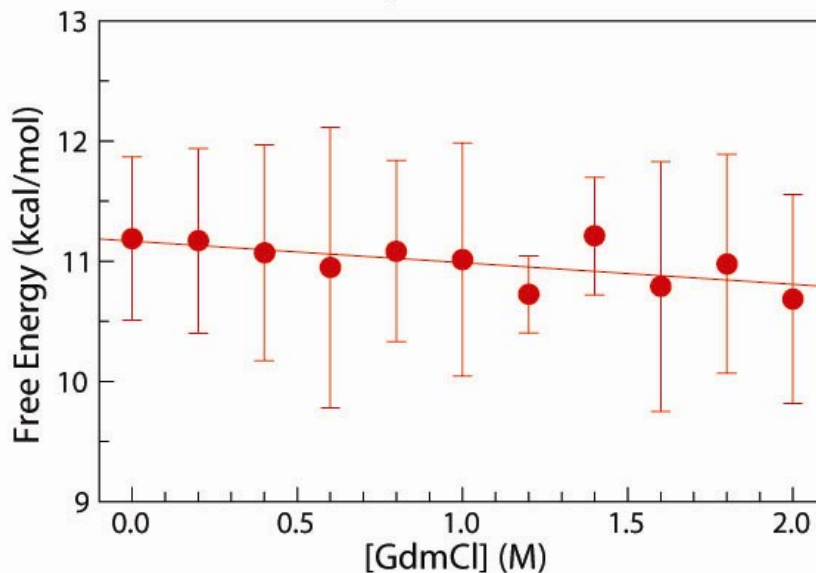
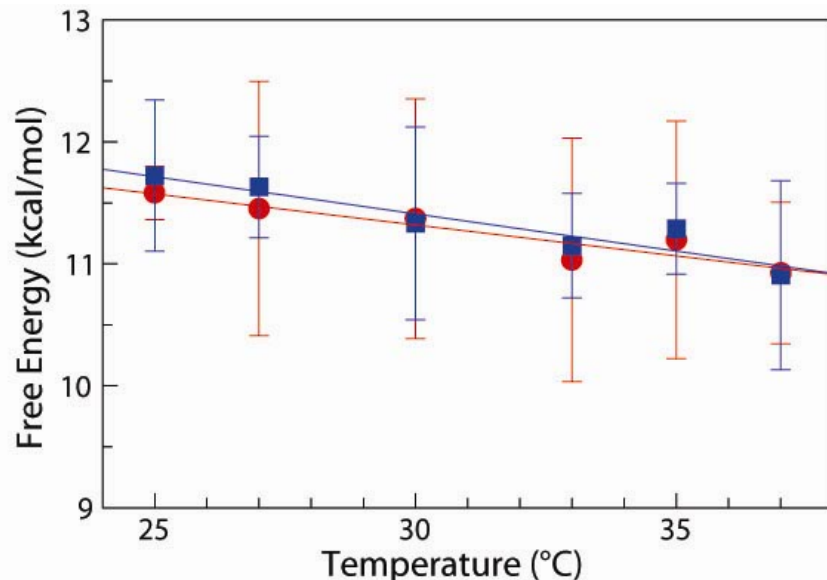


Temperature Dependence of Free Energies



- Titin free energies depends on temperature and denaturant concentration linearly, similar to bulk chemical unfolding.
- Trends consistent for different pulling velocities.

Temperature and Chemical Denaturant Dependent of Unfolding Free Energy Barrier of Titin I27

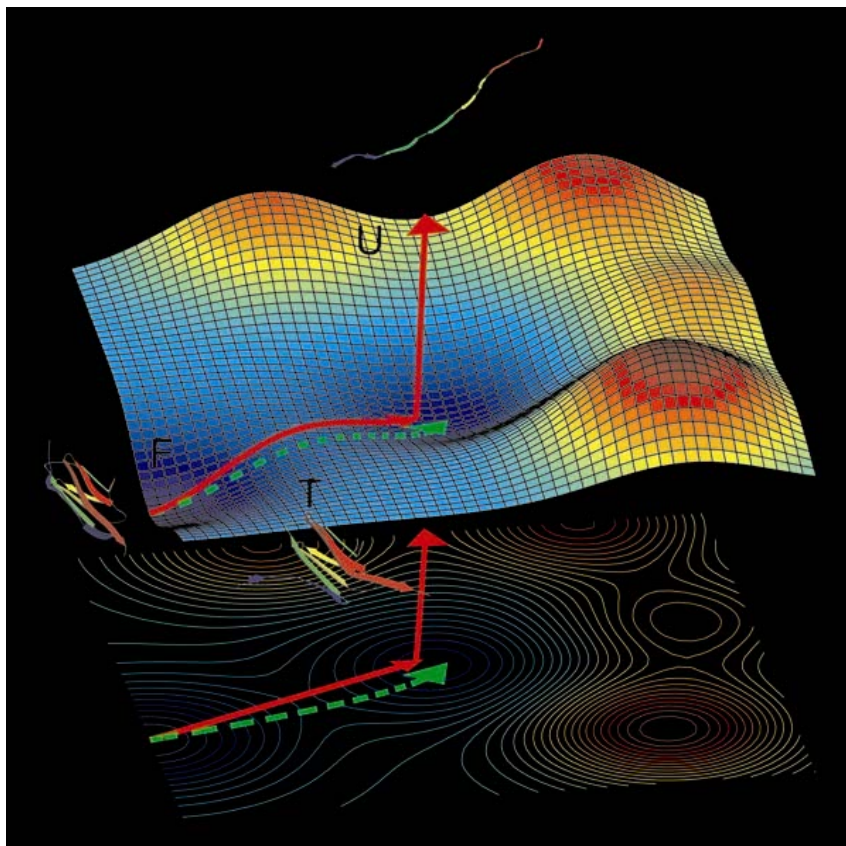


- Trends are consistent with bulk chemical denaturant experiments. $\Delta H^\ddagger=29$ kcal/mol, $\Delta S^\ddagger=0.06$ kcal/mol-T.

Botello, Harris, Sargent, Chen, Lin, and Kiang, **J. Phys. Chem. B** (2009) in press.

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Is End-To-End Distance a Good Reaction Coordinate?



Botello, Harris, Sargent, Chen, Lin, and Kiang, **J. Phys. Chem. B** (2009) in press.

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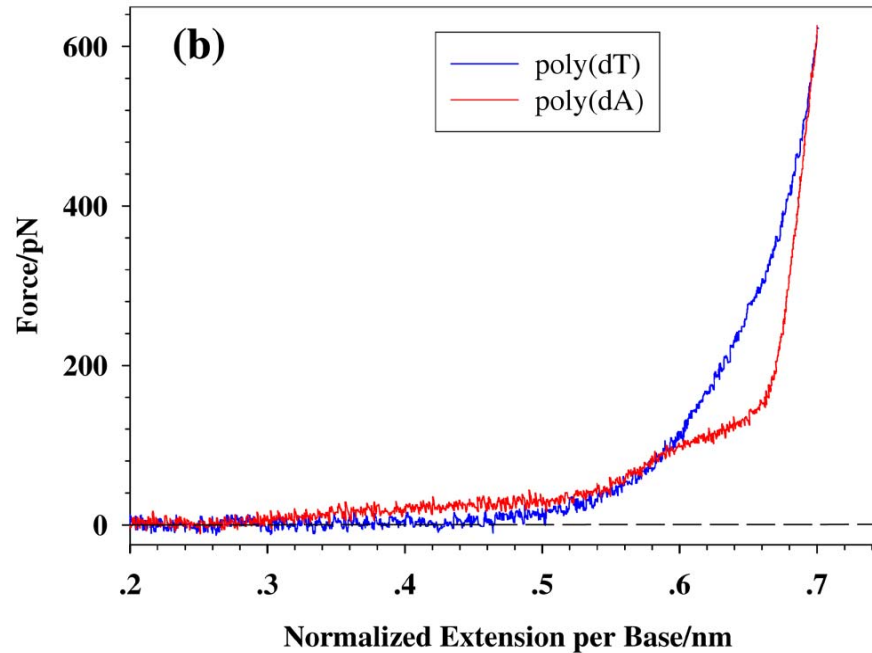
Applications

- Double-stranded DNA melting and overstretching
- Single-stranded DNA unstacking
- RNA folding kinetics
- Protein folding – nanospring
- Protein-protein interactions
- Protein-nucleic acid interactions
- Genome packaging

DNA Melting and Overstretching Transitions

- Conformational and energetic changes of stretched DNA are relevant to biological functions
- During processes such as replication, transcription, and repair, ssDNA is stretched and stabilized by coupling with proteins to serve as an intermediate state
- Ribonucleoprotein complexes in virus (Influenza A, H1N1 virus that causes the “Swine Flu”)
- Implication in microarray analysis
- Base stacking energetics without interacting molecules or melting
- PolydA has two transitions during overstretching transitions
- Elasticity of overstretched ssDNA

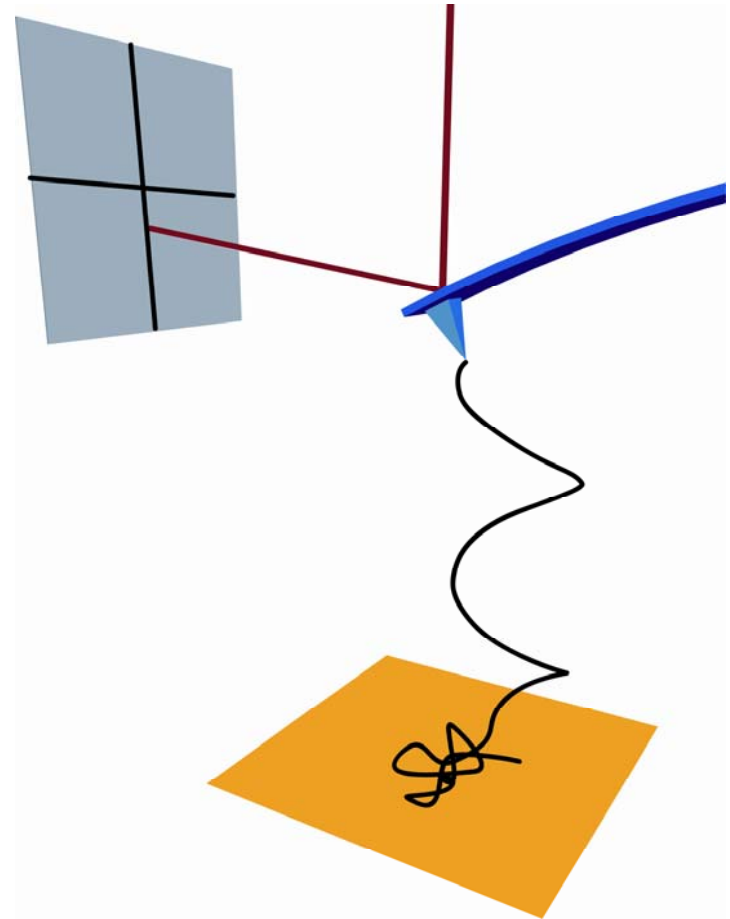
PolydA Exhibits Unique Stacking Behavior



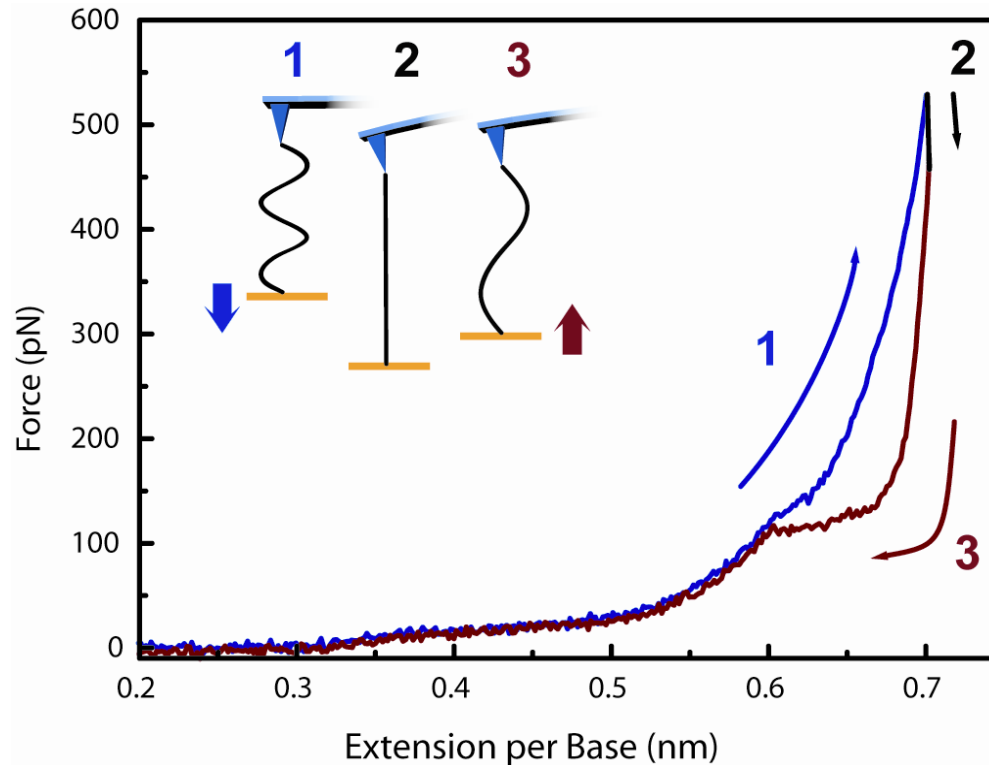
Ke *et. al.*, **Phys. Rev. Lett.**, 99 (2007) 018302.

Pulling Single-Molecules Using Atomic Force Microscopy

- Nanobiology approach to probe biomolecular interactions
- Manipulation and measurements at the single-molecule level
- The end-to-end distance (z) and the force (f) on the molecule were measured

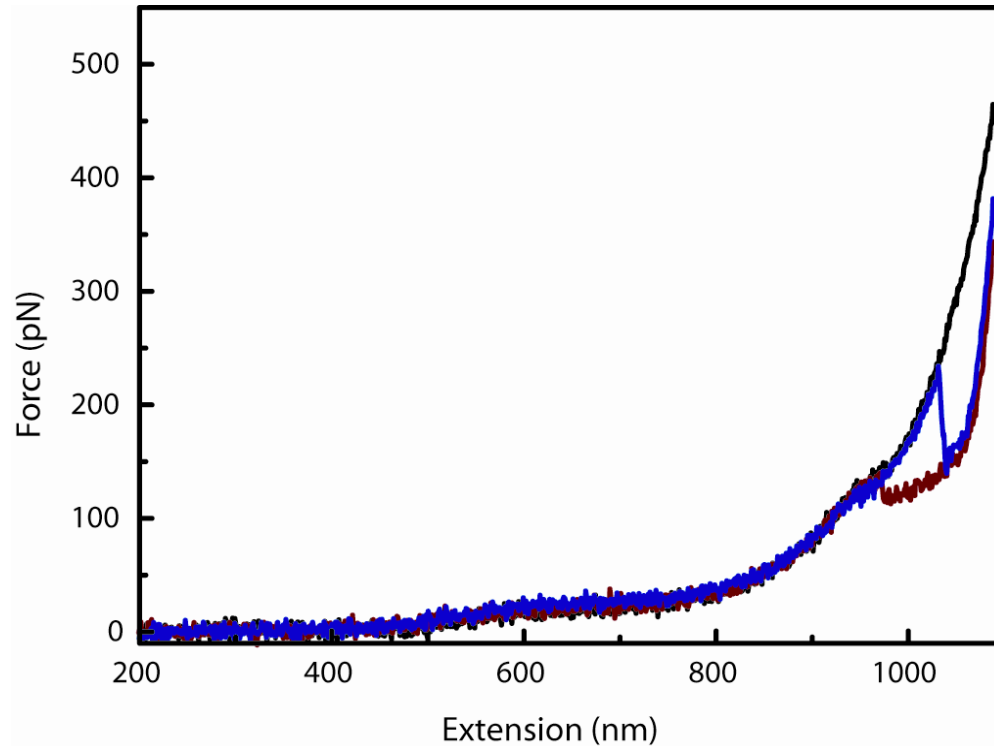


Stacking and Unstacking



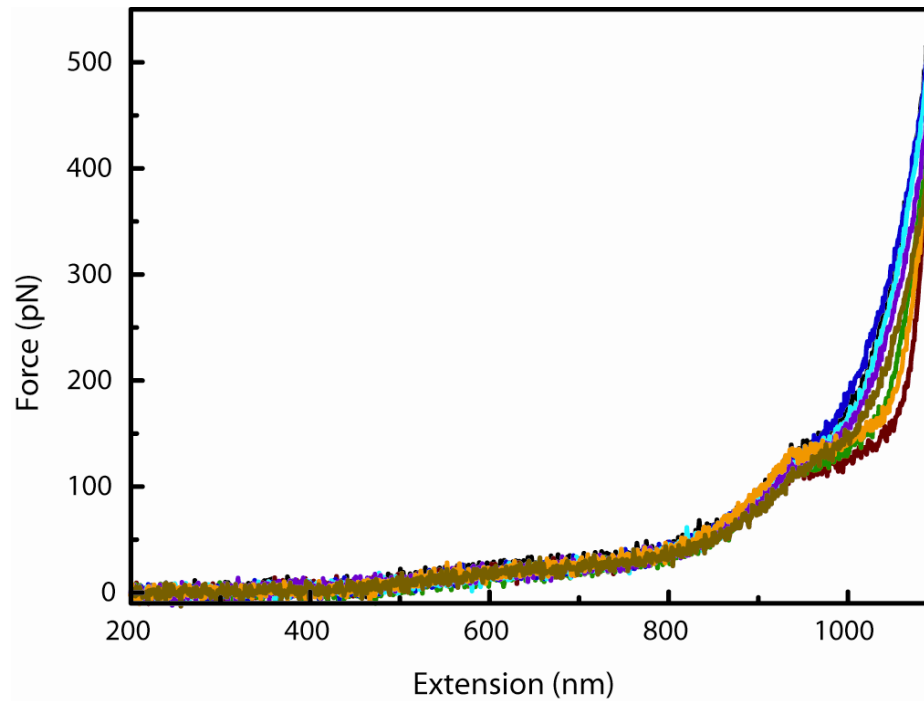
Forward and reverse takes different pathways

Multiple Pathways of PolydA Unstacking



Occasional hopping between two pathways

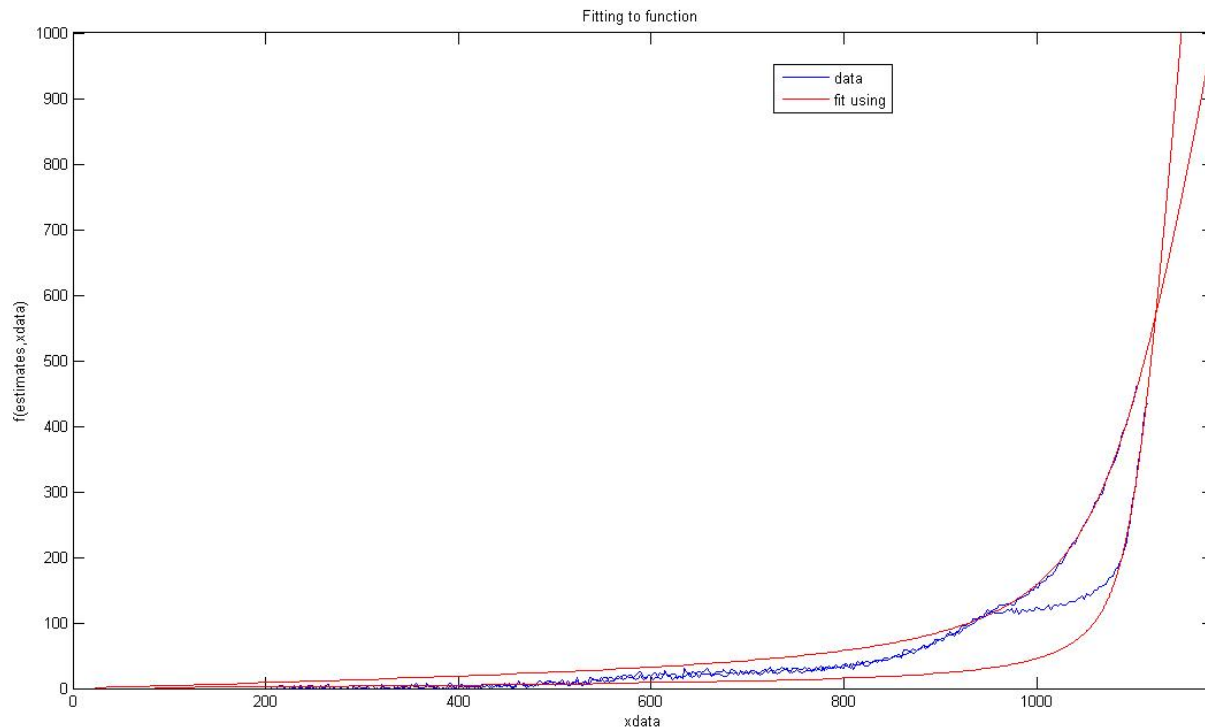
Multiple Stacking Pathways



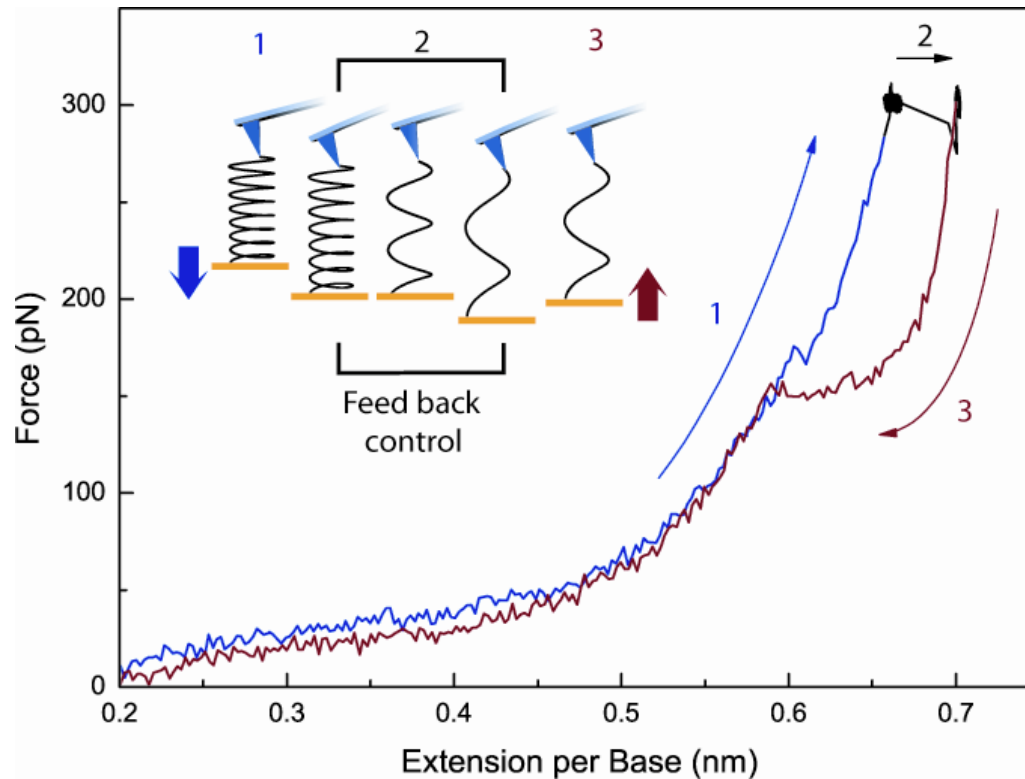
Different persistence length

Two Pathways Intersect

- The two pathways intersect at 600 pN, and the high energy pathway becomes the low energy pathway.

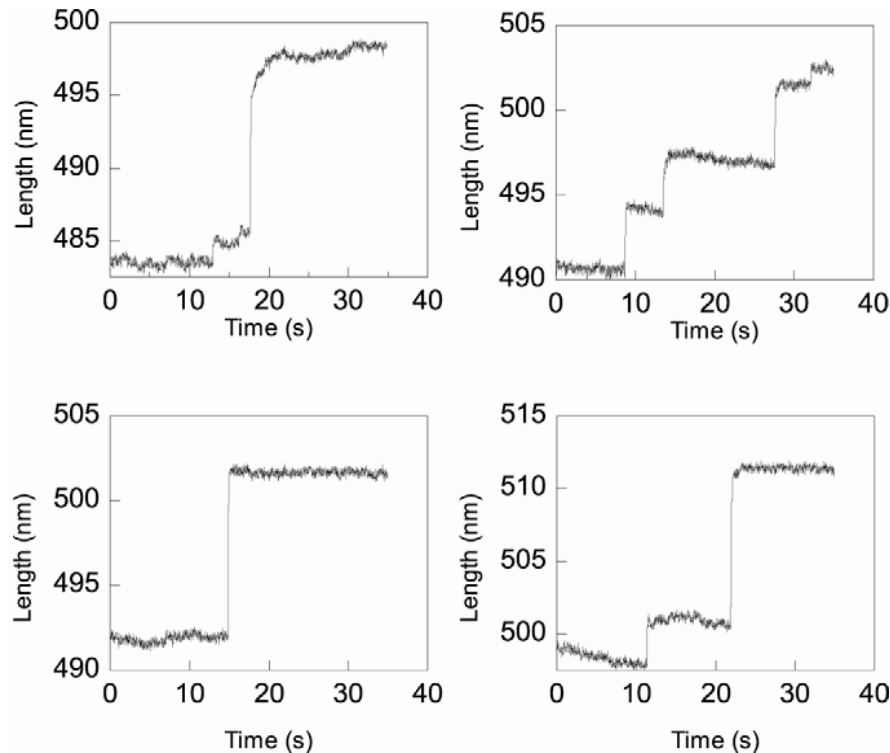


Constant Force Measurements



ssDNA was pulled to 300 pN and kept at constant force

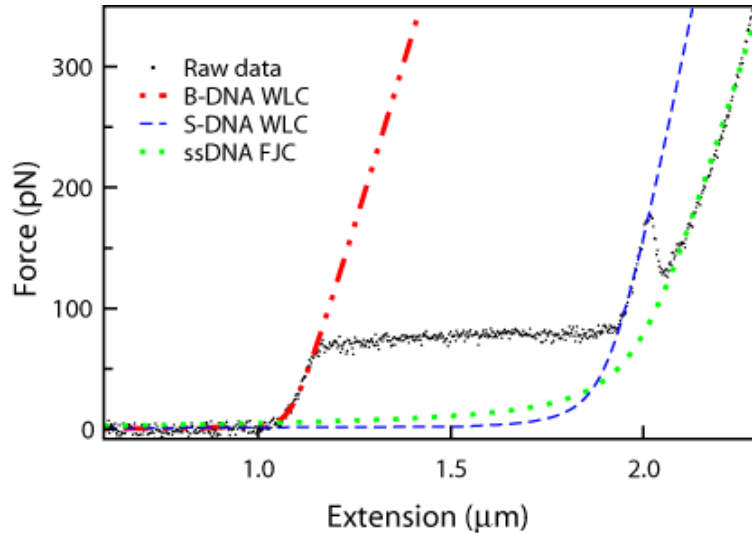
Constant Force Measurements



- Cooperative transitions?
- Ground state configurations?
- dsDNA unzipping metastable states? *Danilowica et.al. PNAS 100, 1694 (2003).*

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Mechanical Melting of Double-Stranded DNA



Extensible Worm-like-chain (WLC)

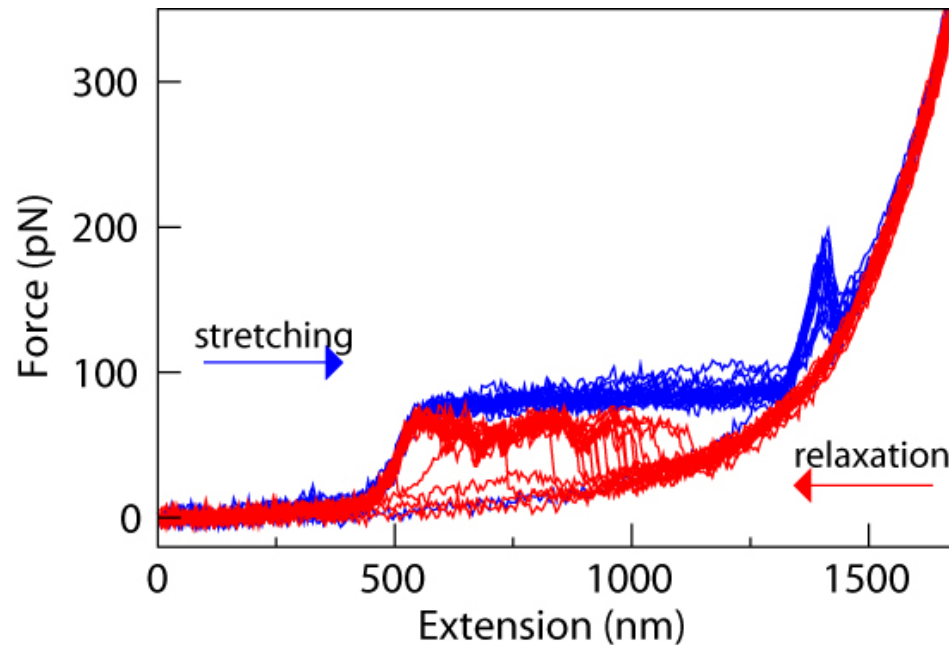
$$x = b_{\text{ds}} \left[1 - \frac{1}{\sqrt{4\beta P_{\text{ds}} F}} + \frac{F}{K_{\text{ds}}} \right]$$

Extensible Freely-Jointed-Chain (FJC)

$$x = b_{\text{ss}} \left[\coth(2\beta P_{\text{ss}} F) - \frac{1}{2\beta P_{\text{ss}} F} \right] \left[1 + \frac{F}{K_{\text{ss}}} \right]$$

	P persistence Length (nm)	b contour length (nm)	K stretch modulus (pN)
<i>B-DNA</i>	50	1100	1200
<i>S-DNA</i>	10	2000	3200
<i>ssDNA</i>	0.75	2000	2200

B-S Transition of *dsDNA*



- Repeated stretch/relaxation cycle reproducible
- Significant hysteresis during ssDNA relaxation cycle

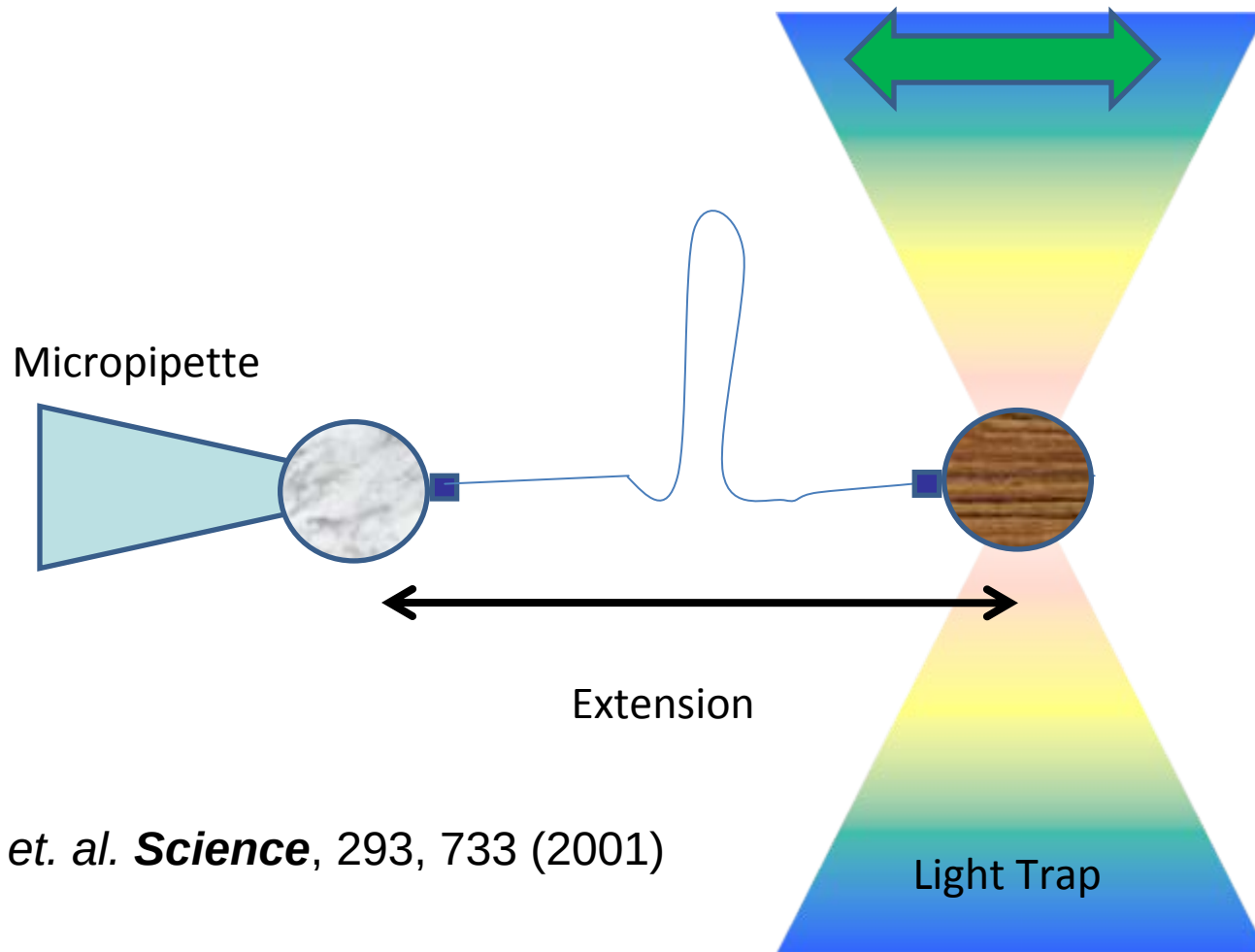
Calderon, Harris, Kiang, and Cox, J. Phys. Chem. B 113, 138 (2009).

Calderon, Harris, Kiang, and Cox, J. Mol. Recognit., (2009) in press.

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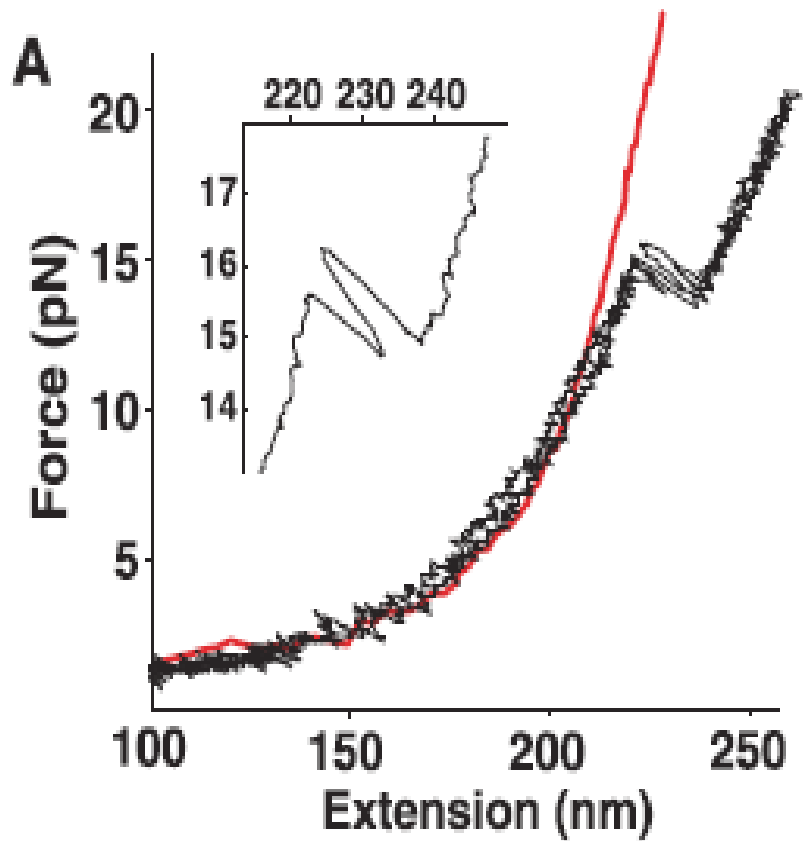
Force Curve Setup

Move Trap and Measure Force

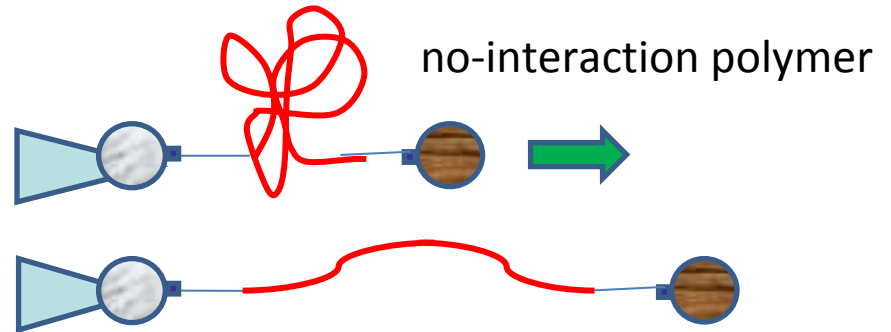


Lipardt et. al. **Science**, 293, 733 (2001)

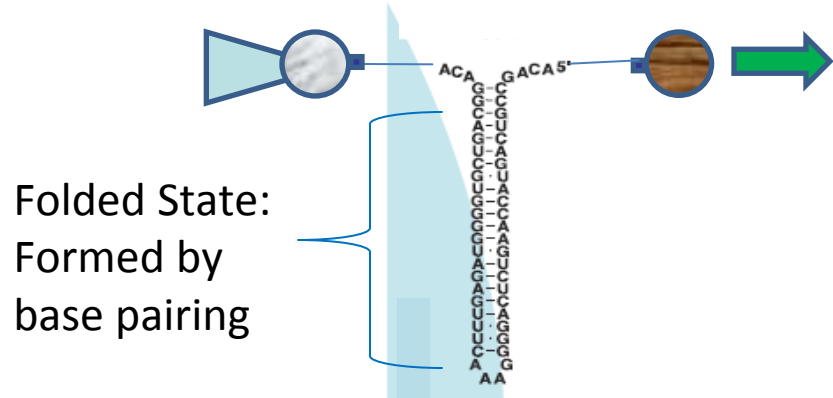
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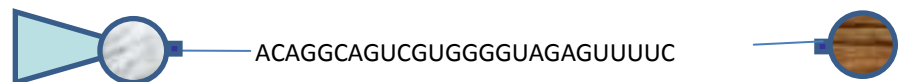
Red Line (Satisfies WLC):



Black Line (Hairpin RNA):



Unfolded State:
Hydrogen Bonds Broken by Force



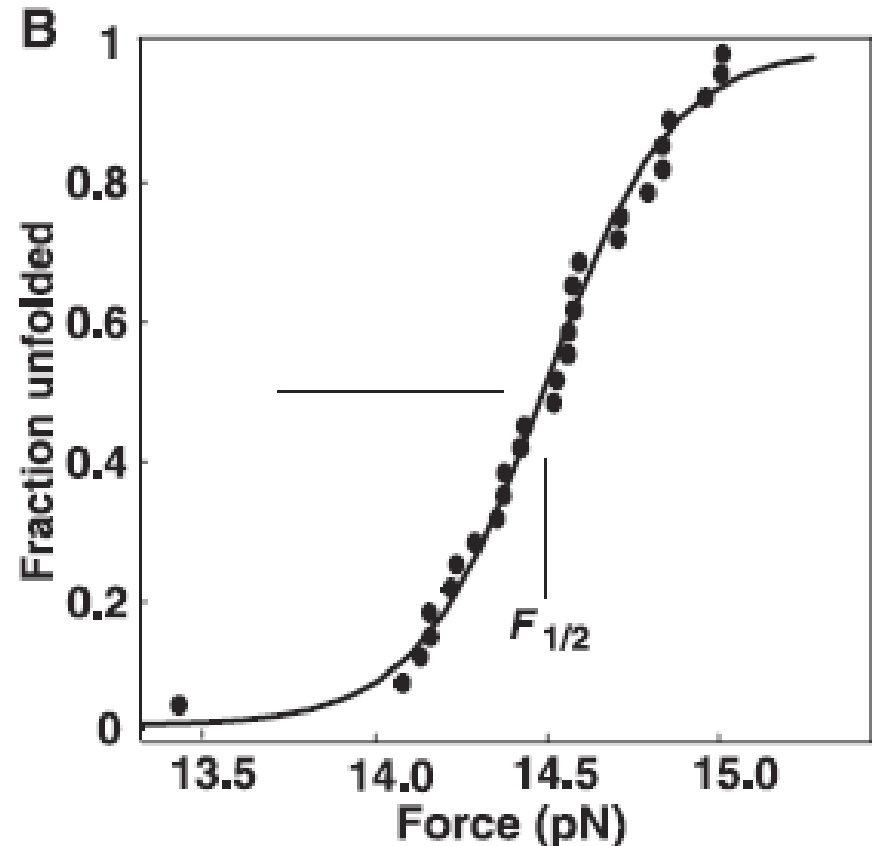
Equilibrium Approaches

- By repeated folding/unfolding, one can get a graph of fraction unfolded RNA versus force and fit into a two-state equation:

$$p(E) = 1/(1 + e^{E/k_B T})$$

- The free energy of unfolding of RNA

$$\Delta G = 157 \pm 20 \text{ kJ / mole}$$

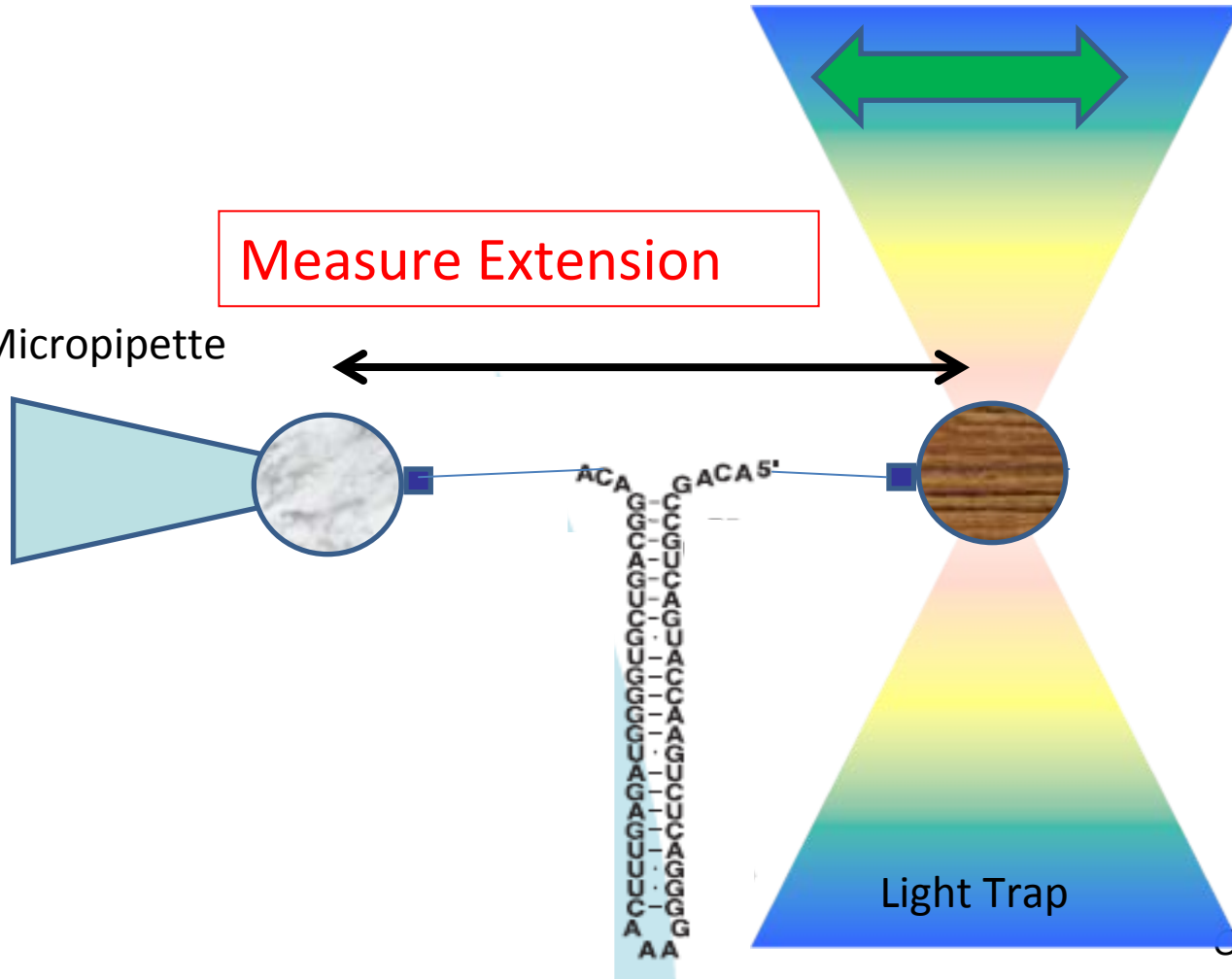


Constant Force Setup

Maintain force by moving the trap

Measure Extension

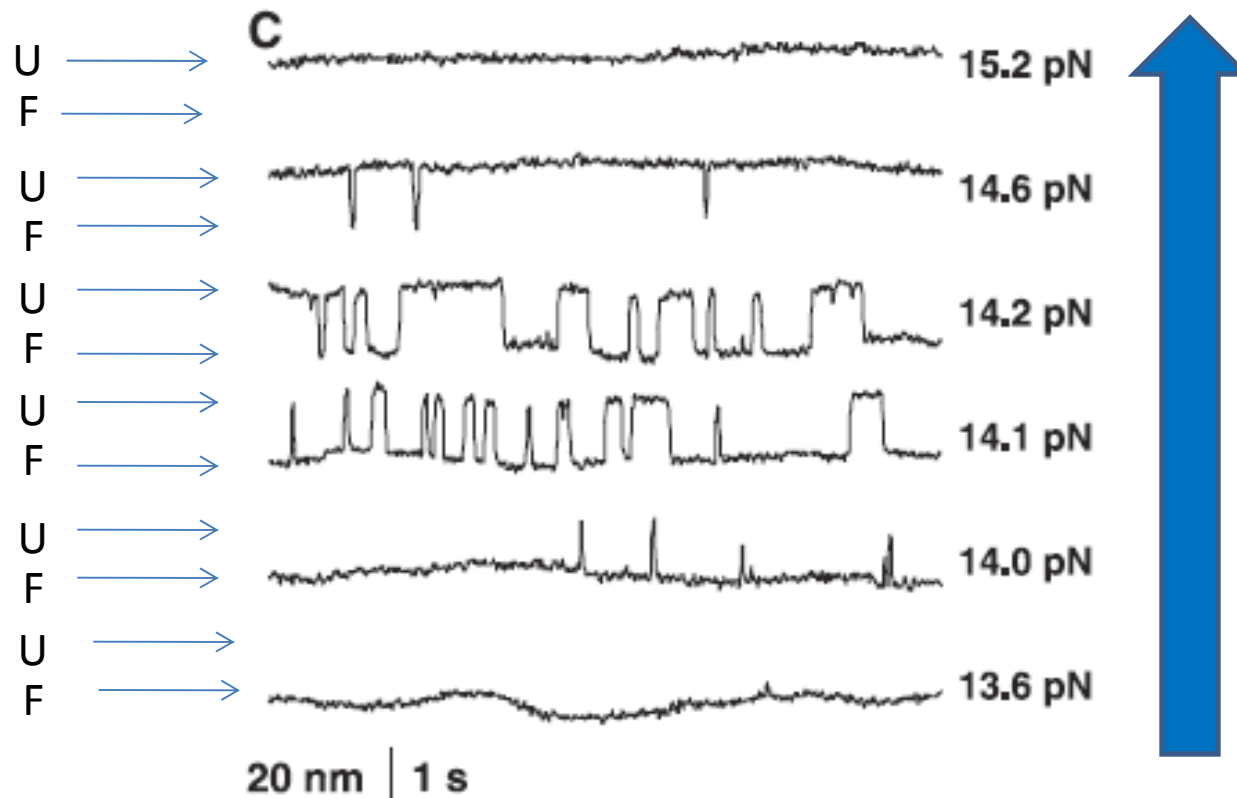
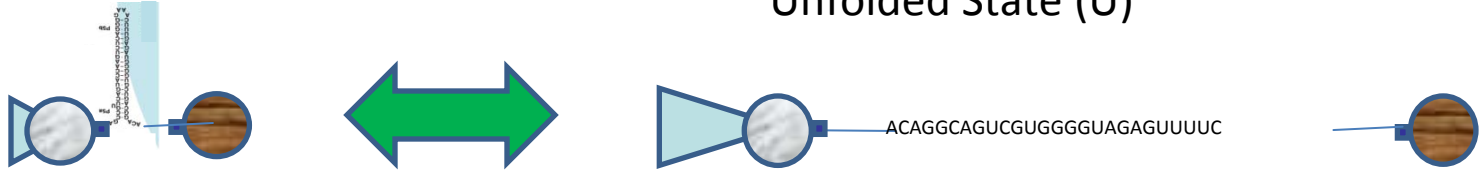
Micropipette



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Folded State (F)

Unfolded State (U)



High force
favors open
state

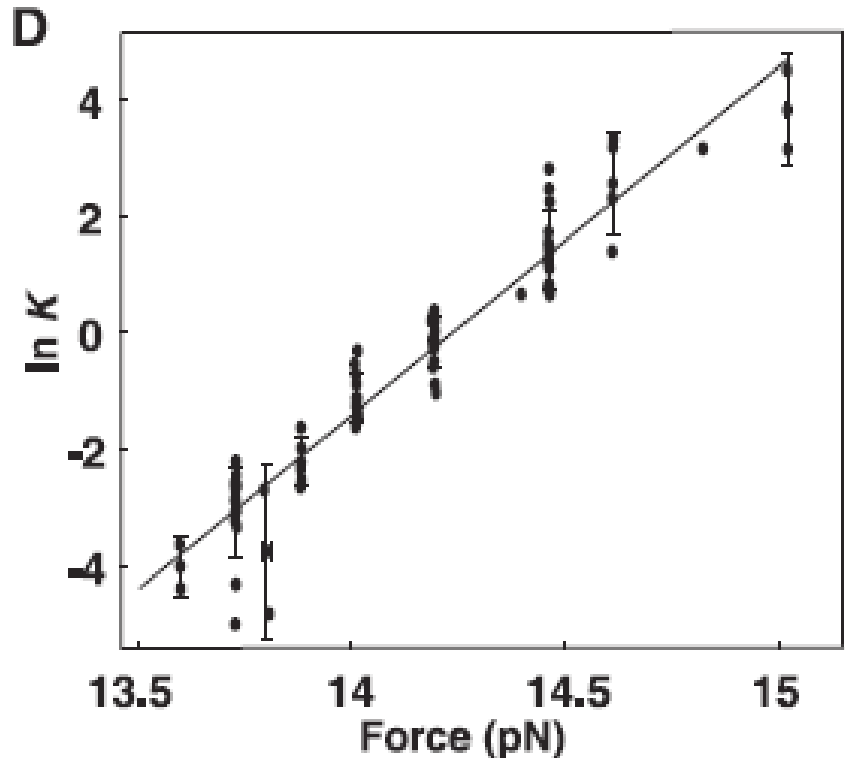
Free Energy Estimate

- The ratio of the average life time results in K . By plotting $\ln K$ versus force, we can get the free energy between the unfold and folded state using:

$$\Delta G = \Delta G^o + RT \ln K$$

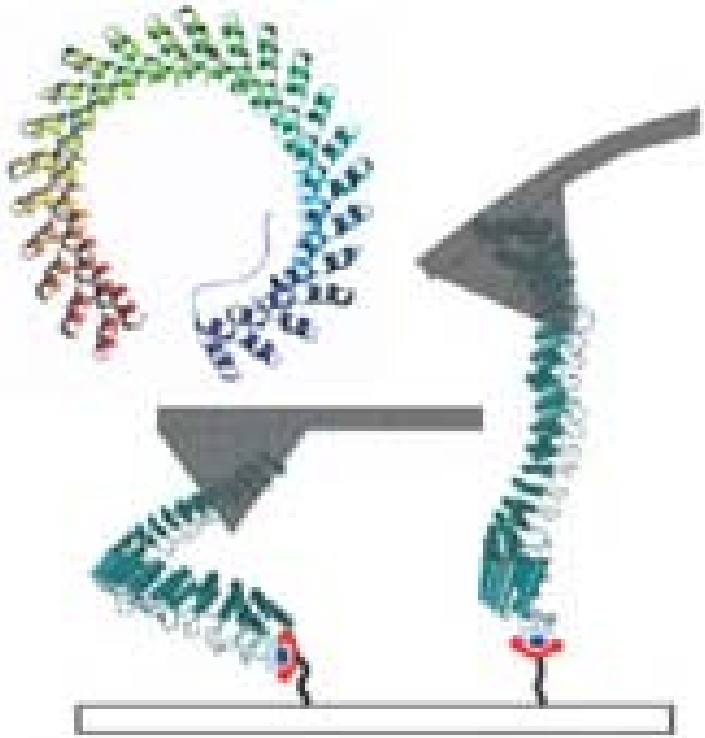
They obtained

$\Delta G = 156 \pm 8 \text{ kJ / mole}$
consistence with bulk studies.



Protein Nanosprings

Linear Elasticity



His-tagged ankyrin fragment
bound to NTA (red handles)

Unfolding Ankyrin Repeats

