

# Viral DNA as a model for coil to globule transition

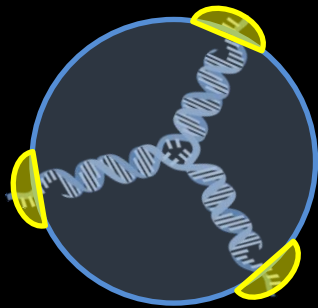
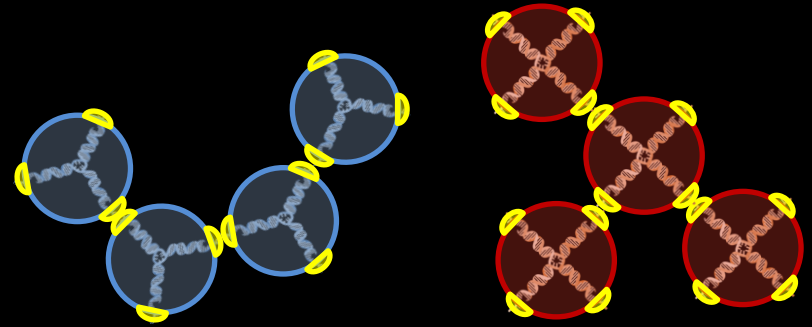


Marina Rossi

Lab. of complex fluids and molecular biophysics  
LITA (Segrate)

# DNA AS A MOLECULAR MODEL

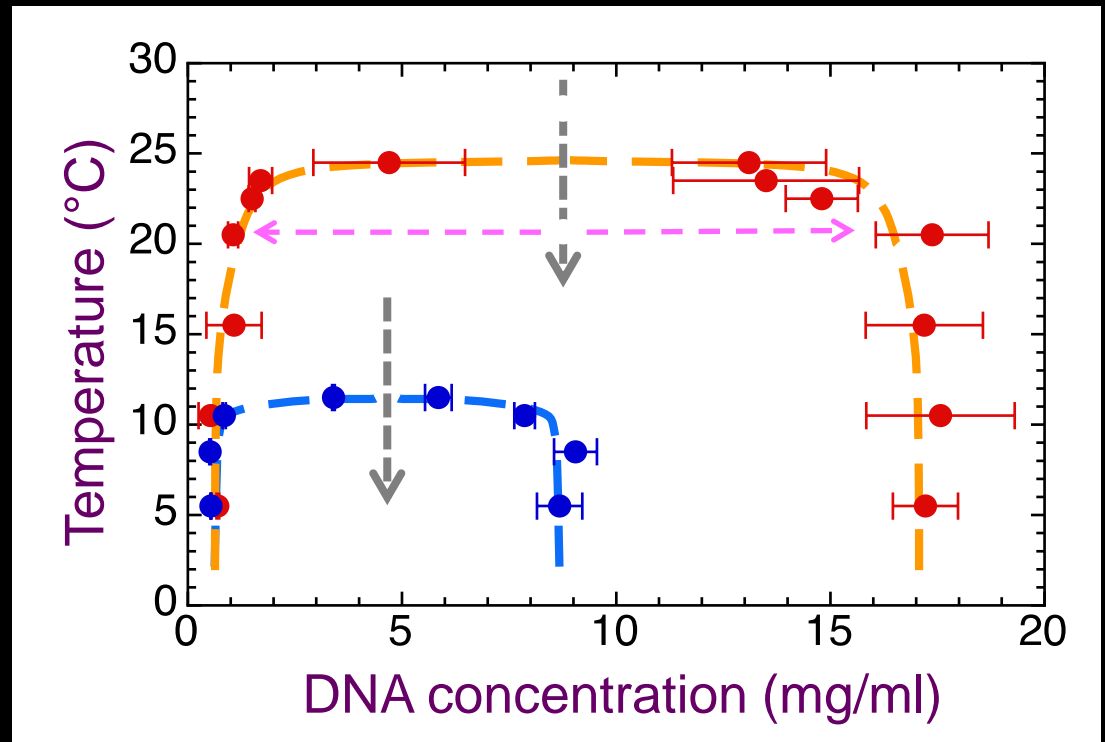
DNA for modeling particles at finite valence



3-valence



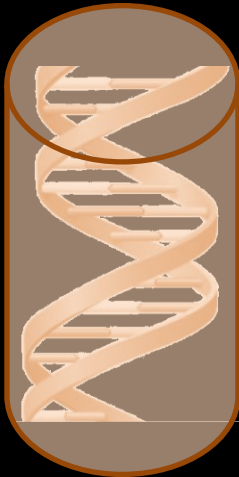
4-valence





( last year )

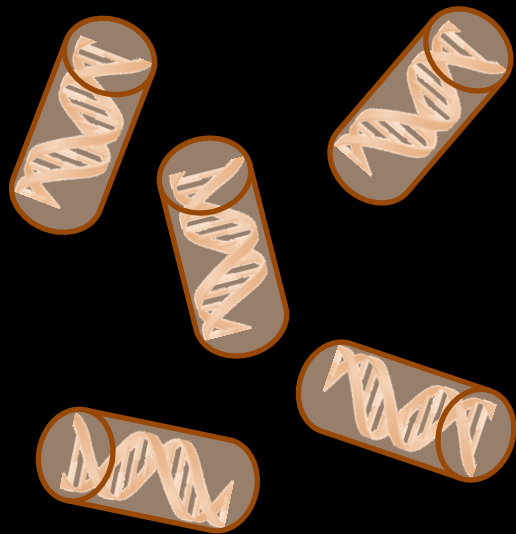
# Liquid crystalline phases of DNA



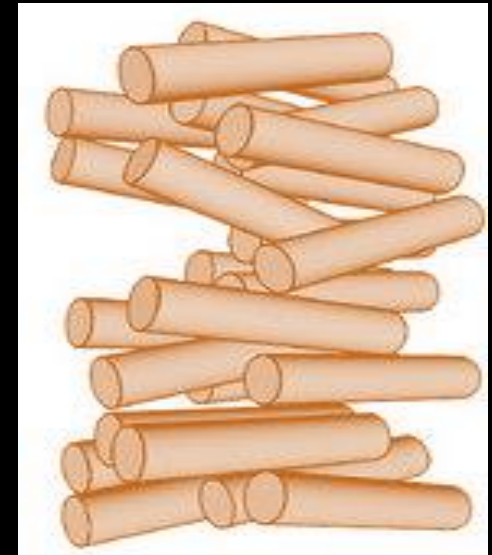


( last year )

## Liquid crystalline phases of DNA



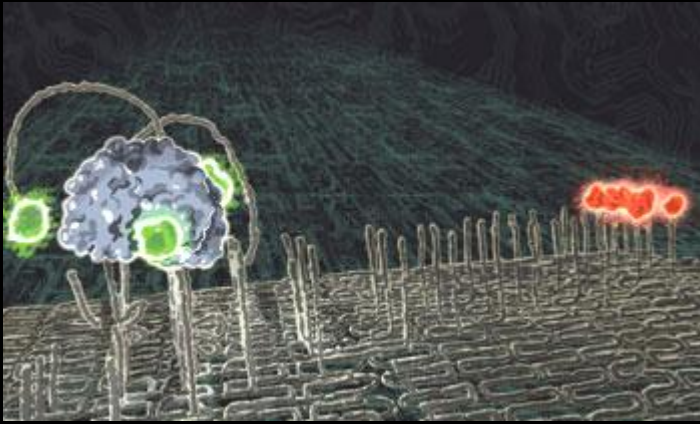
high concentration



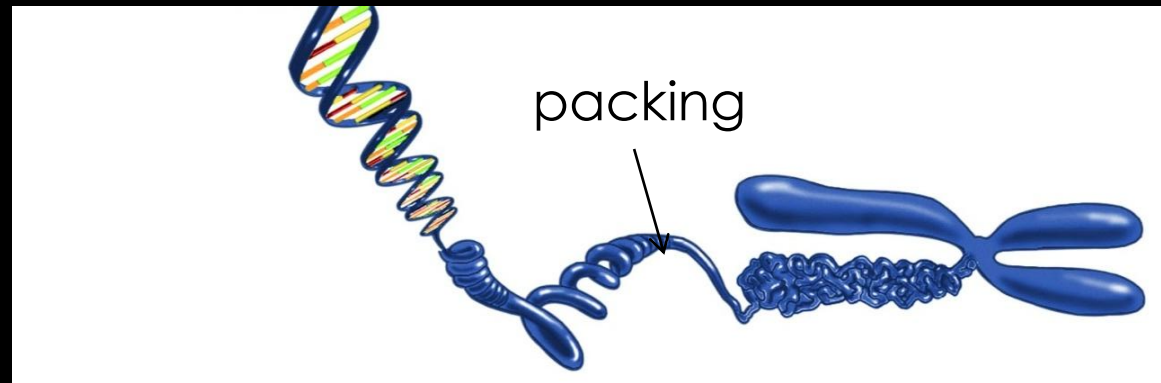
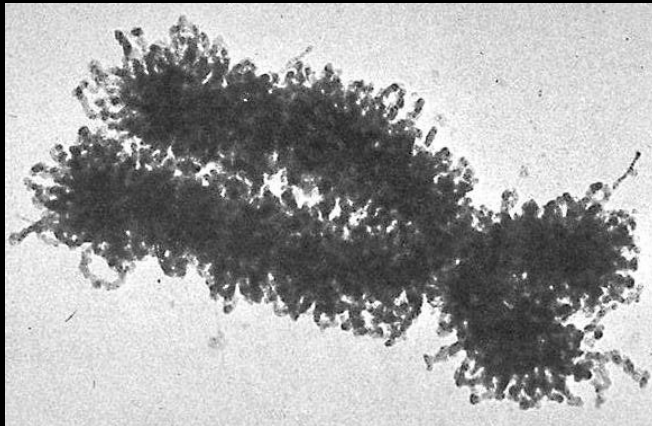
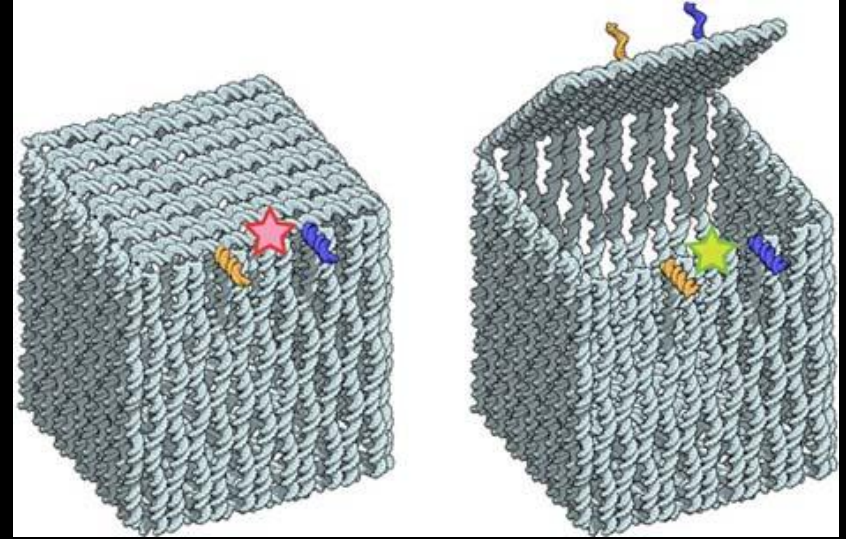
Properties of DNA  $\longrightarrow$  properties of the LC phase

M. Rossi *et al.* PRL **110** 107801 (2013)

# Applications: DNA nanotechnologies



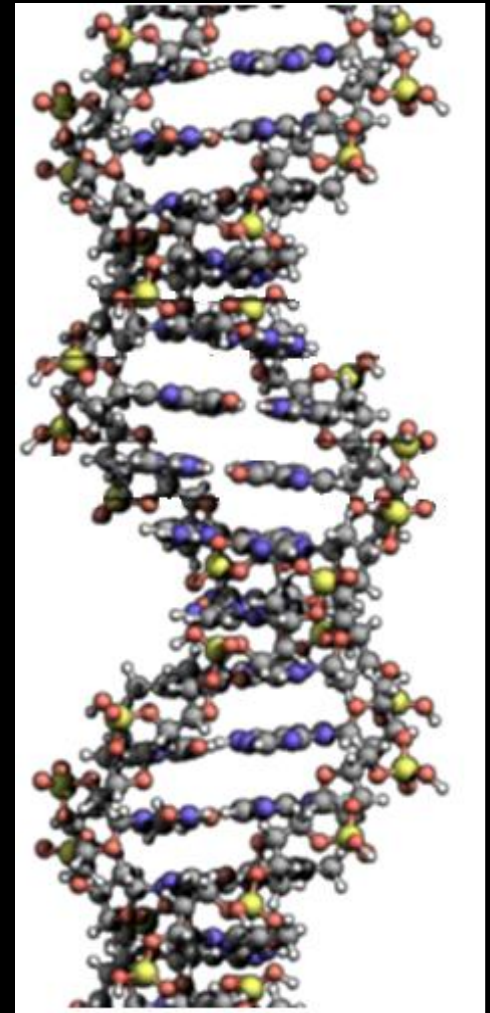
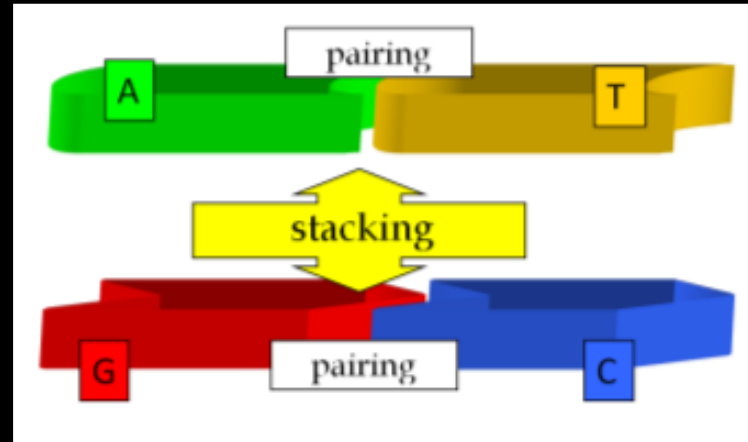
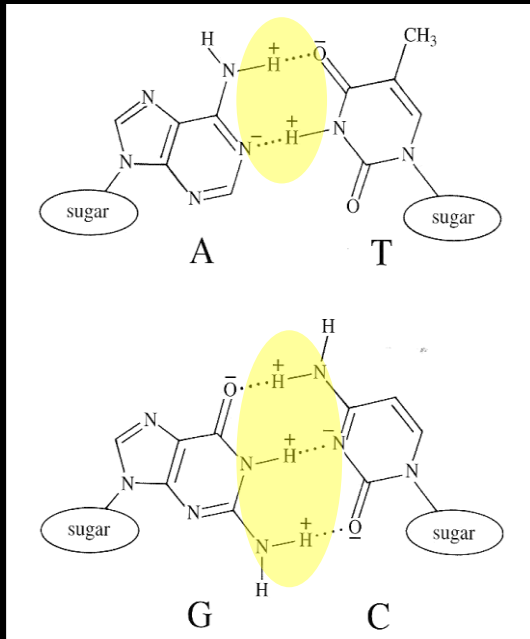
Chromosomes



Human Chromosome 12 -electron microscopy, E. Du Praw

# Why DNA?

- Well known structure
- Selective interactions

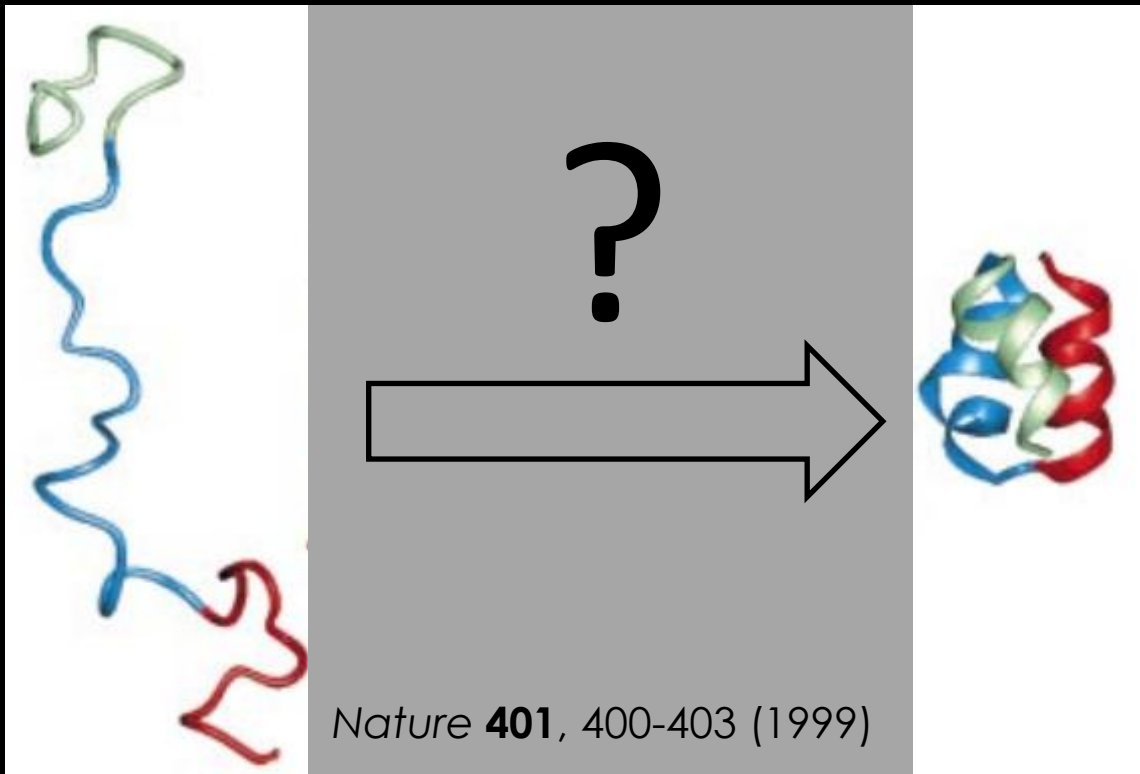


- Tunable interactions  
(temperature, ionic strength, concentration,..)

# CHAPTER 2

(Today)

DNA to study coil to globule transition



Protein folding

Denatured state  
Not functional



native state  
Functional

Protein:

Non specific interactions

- hydrogen bonds
- hydrophobic
- electrostatic
- steric
- Van der Waals

Sequence  $\longrightarrow$  unique3D folded structure



Protein:

Non specific interactions

- hydrogen bonds
- hydrophobic
- electrostatic
- steric

- Van der Waals

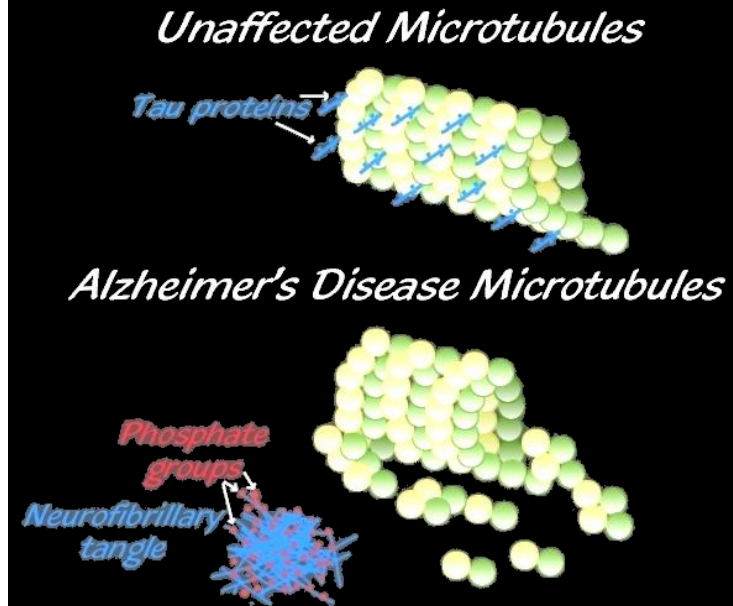
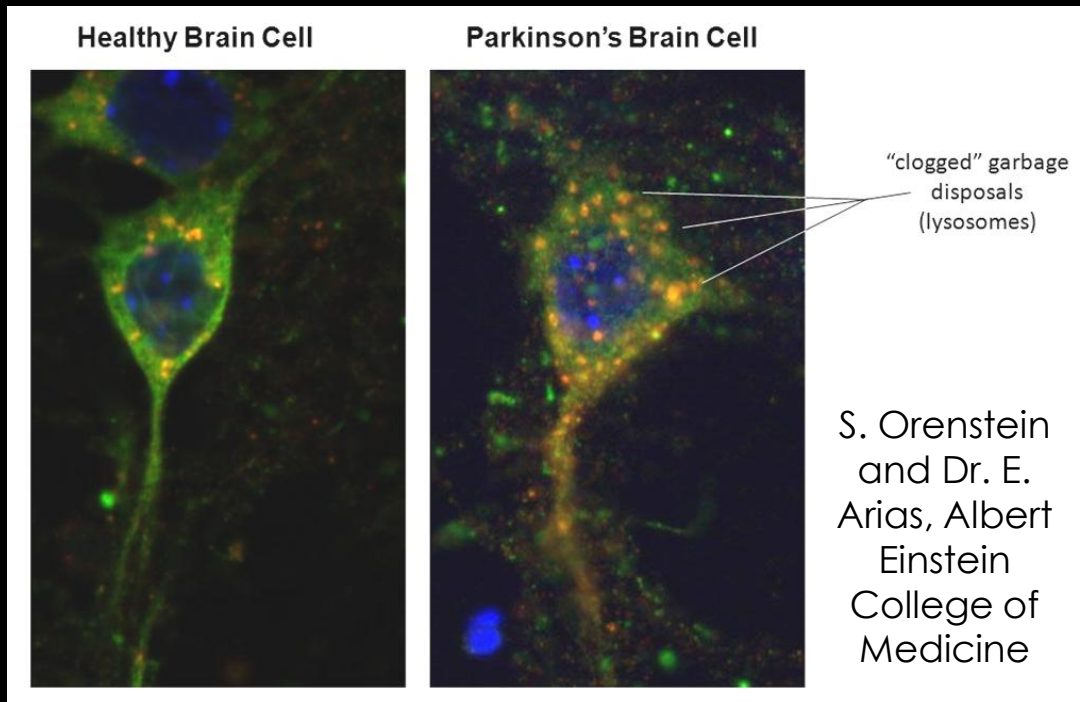
Sequence  $\longrightarrow$  unique 3D folded structure

Unresolved questions

- how does the native structure result from the **interatomic forces** ?
- how can the native structure be predicted from the **sequence** ?
- how can a protein fold so **fast** ?

# Why folding process?

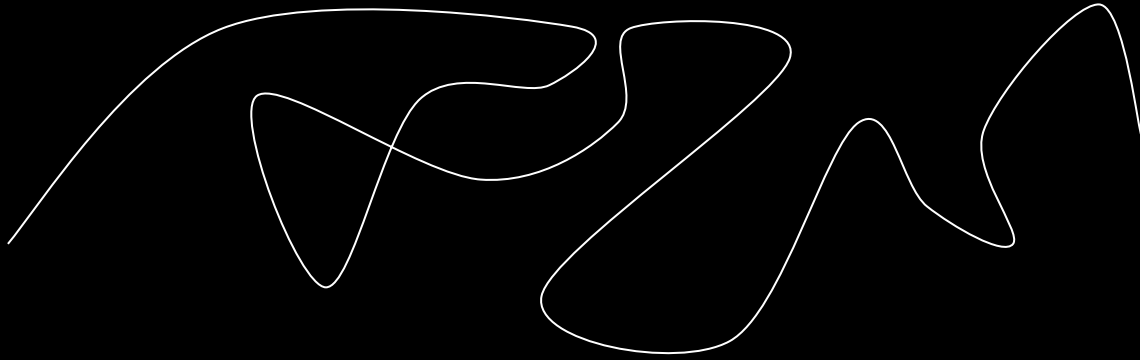
- change of native structure
  - Tau protein/  $\alpha$ -synuclein :  
Alzheimer's and Parkinson's diseases



- Protein inhibition
  - HIV-1-protease

# DNA: a model for proteins

Viral sDNA: a heteropolymer



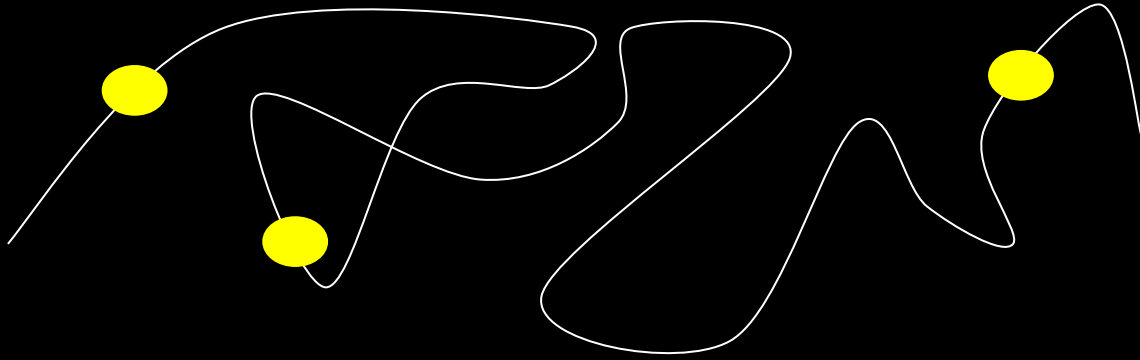
AFM image of filamentous phage



Mol. Biol. **13**, 51

# DNA: a model for proteins

Viral sDNA: a heteropolymer



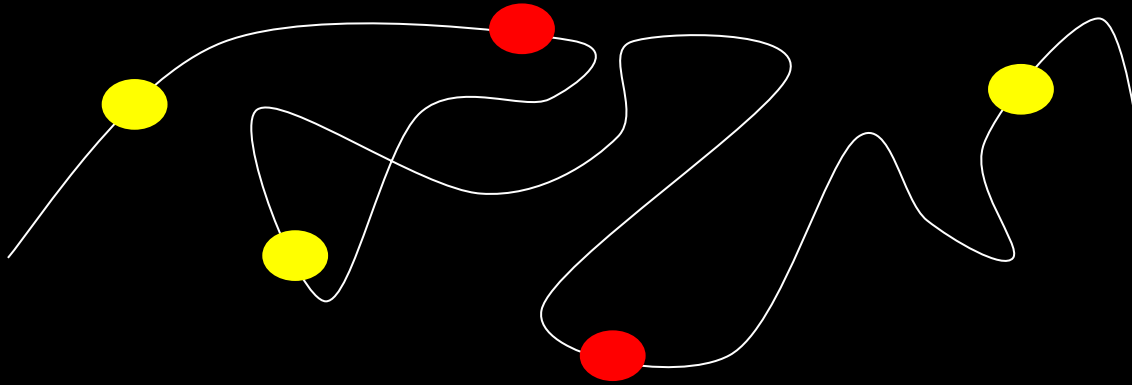
AFM image of filamentous phage



Mol. Biol. **13**, 51

# DNA: a model for proteins

Viral sDNA: a heteropolymer



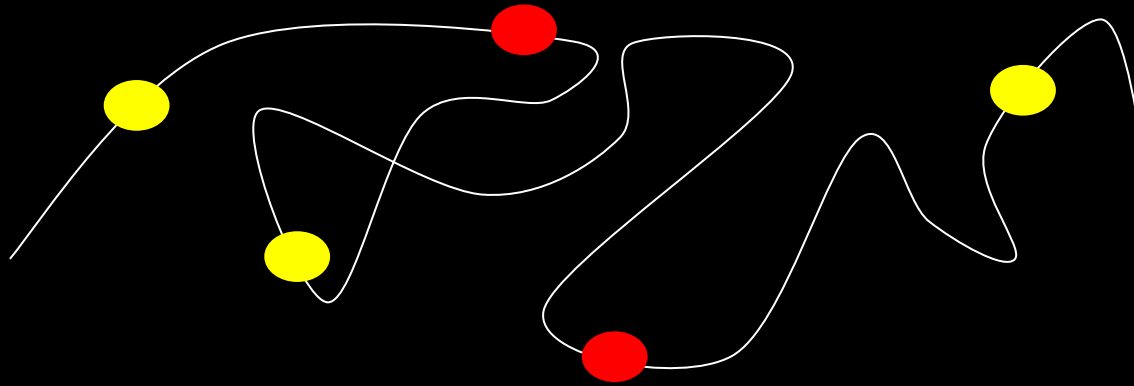
AFM image of filamentous phage



Mol. Biol. **13**, 51

# DNA: a model for proteins

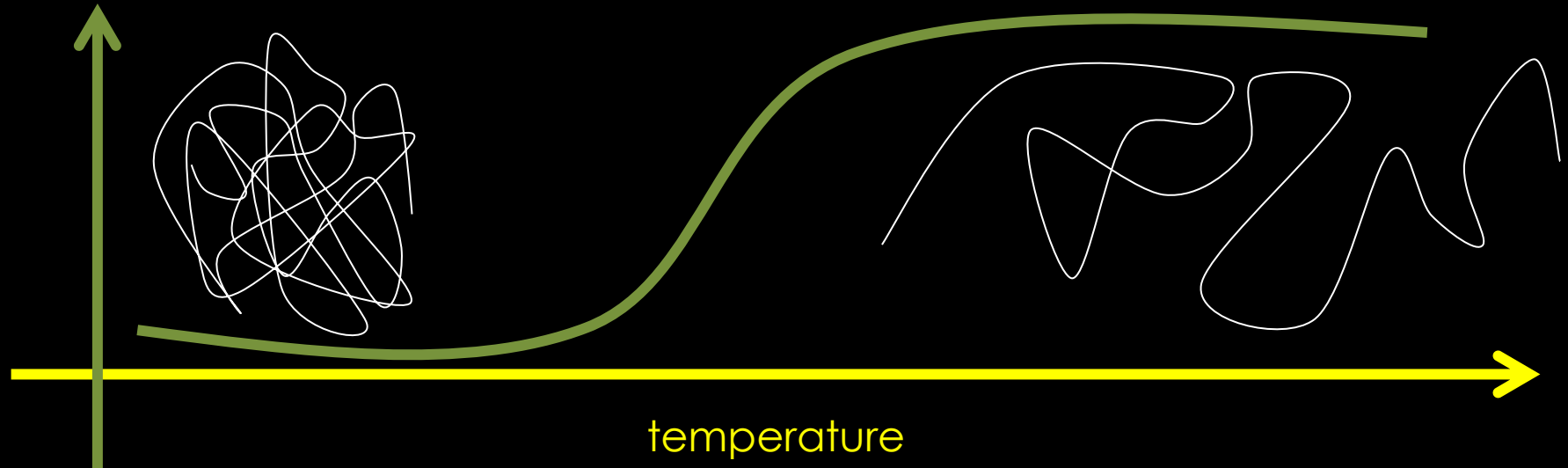
Viral sDNA: a heteropolymer



AFM image of filamentous phage  
Mol. Biol. **13**, 51



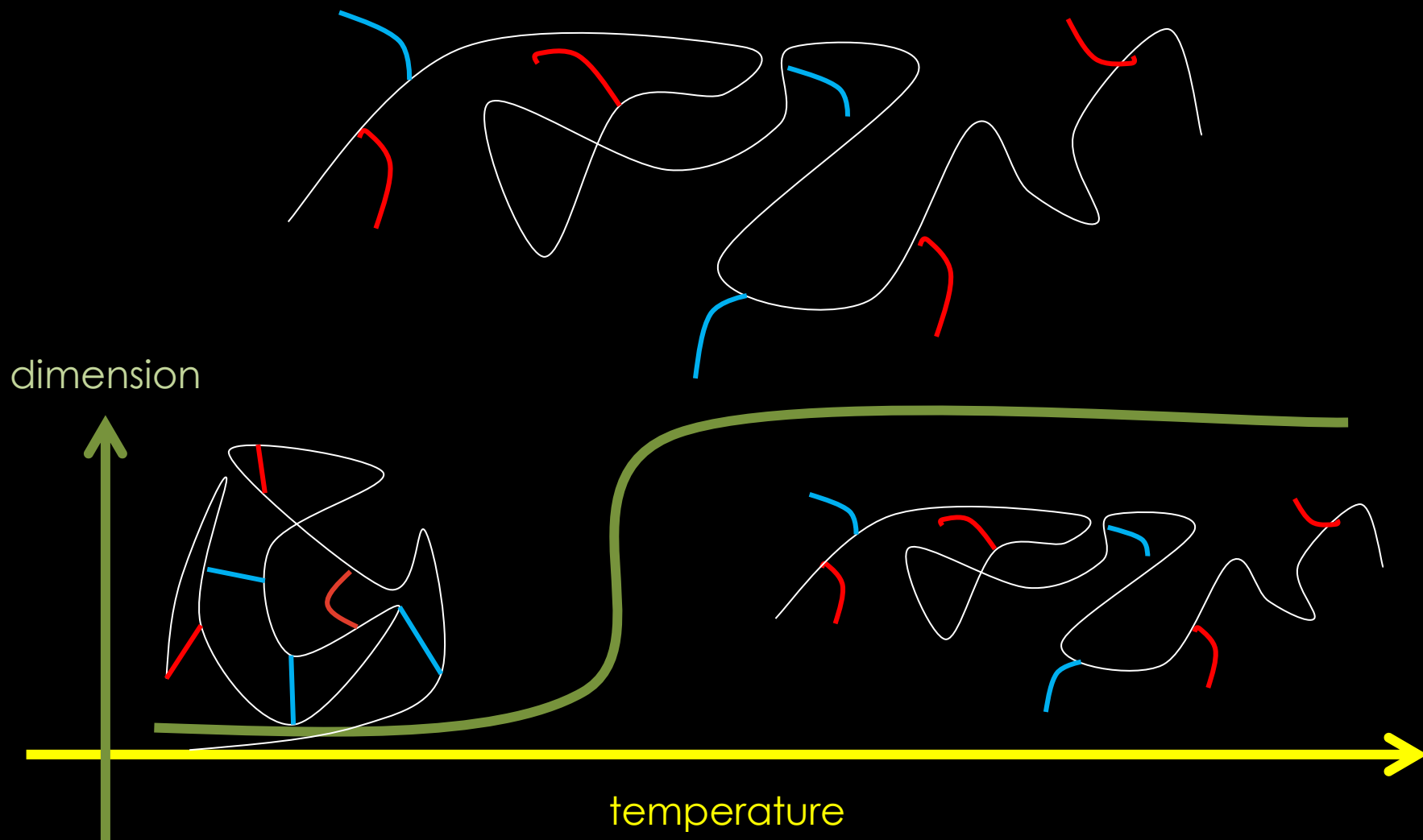
dimension



# M13mp18 – 5000 bases out of 7000

AATGCTACTACTATTAGTAGAATTGATGCCACCTTTTCAGCTCGCGCCCCAAATGAAAATATAGCTAAACAGGTTATTGACCAATTGCGAAATGTATCTA  
ATGGTCAAACATAAATCTACTCGTTCGCAGAATTGGGAATCAACTGTTATATGGAATGAAACTTCCAGACACCGTACTTTAGTTGCATATTTAAACATGTT  
GAGCTACAGCATTATATTCAGCAATTAAGC **CTAAGCCATC** CGCAAAAATGACCTCTATCAAAGGAGCAATTAAGGTACTCTCTAATCCTGACCT  
GTTGGAGTTTGCTCCGGTCTGGTTCGCTTTGAAGCTCGAATTAACCGCGATATTTGAAGTCTTCGGGCTTCTCTAATCTTTTATGCAATCCGCTTG  
CTTCTGACTATAATAGTCAGGGTAAAGACCTGATTTTGAATATGGTCATCTCGTTTCTGAACTGTTAAAGCATTGAGGGGGATTCAATGAATATTTATG  
ACGATTCCGCAGTATTGGACGCTATCCAGTCTAAACATTTACTATTACCCCTCTGGCAAACCTCTTTTGCAAAAGCCCTCTCGCTATTTGGTTTTATCG  
TCGTCTGGTAAACGAGGGTATGATAGTGTGCTCTACTATGCCTCGTAATTCCTTTGGCGTATGTATCTGC **ATTAGITGAA** GTGGTATTCCTAAATCTCA  
ACTGATGAATCTTTCTACCTGAATAATGTTGTTCCGTTAGTTCG **TTTTATTA** CGTAGATTTTCTTCCCAACGTCTGACTGGTATAATGAGCCAGTCTTAAA  
ATCGCATAAGGTAATTCACAATGATTAAGTTGAAATTAACCATCTCAAGCCCAATTACTACTCGTTCGGTGTCTCGTCAGGGGCAAGCCATTATCA  
CTGAATGAGCAG **ATTCAACTAAT** CTTTGTACGTTGATTGGGTAATGAATATCCGGTCTTGTCAAGATTACTCTTGATGAAGGTCAGCCAGCCTATGCG  
CCTGGTCTGTACACCGTTCATCTGTCTCTTTCAAAGTTGGTAGTTCGGTCCCTTATGATTGACCGTCTGCGCCTCGTCCGGCTAAGTAACATGGAGC  
AGGTCGCGGATTCGACACAATTTATCAGGCGATGATACAAATCTCCGTTGTACTTTGTTTCGCGCTTGGTATAATCGCTGGGG **GTCAAAG** ATGAGTGT  
TTAGTGTATTCTTTGCCTCTTTGTTTTAGGTTGGTGCCTC **CTTTGAC** GTAGTGGCATTACGTATTTACCCGTTAATGGAACTTCCTCATGAAAAAGTCTT  
AGTCTCAAAGCCTCTGTAGCCGTTGCTACCCTCGTCCGATGCTGTCTTTGCTGCTGAGGGTGACGATCCCGCAAAGCGGCCTTTAACTCCCT  
GCAAGCCTCAGCGACCGAATATATCGGTTATGCGTGGGCGATGGTGTGTCATTGTCGGCGCAACTATCGGTATCAAGCTGTTAAGAAATTCACCT  
CGAAAGCAAGCTGATAAACCGATACAATAAAGGCTCCTTTGGAGCCTTTTTTTGGAGATTTCAACGTGAAAAAATTATATTCCGAATTCCTTAGTTG  
TTCCTTCTATTCTCACTCCGCTGAAACTGTTGAAAGTTGTTAGCAAATCCCATAC **AGAAAATTA** TACTAACGCTGGAAAGACGACAAAACCTTAG  
ATCGTTACG **CTAACTATG** AGGGCTGTCTGTGGATGCTACAGGCGTTGTAGTTTGTACTGGTGACGAAACTCAGTGTACGGTACATGGGTCCCTATTGG  
GCTTGTATCCCTGAAAATGAGGGTGGTGGCTCTGAGGGTGGCGGTTCTGAGGGTGGCGGTTCTGAGGGTGGCGGTAATAACCTCCTGAGTAC  
GGTGATACACCTATCCGGGCTATACTTATATCA **ACCCTC** TGACGGCACTTATCCGCCTGGTACTGAGCAAACCCCGCTAATCCTAATCCTTCTCT  
GAGGAGTCTCAGCCTCTAATACTTTTATGTTTCAAGAATAATAGGTTCCGAAATAGGCAGGGGGCATTAACTGTTTATACGGGCACTGTTACTCAAGGC  
ACTGACCCCGTAAAACCTATTACCAGTACACTCCTGTATCATCAAAGCCATGTATGACGCTTACTGGAACGGTAAATTCAGAGACTGCGCTTTCCAT  
TCTGGCTTAATGAGGATTTATTTGTTTGAATATCAAGGCCAATCGTCTGACCTGCCTCAACCTCCTGTCAATGCTGGCGGGCGGCTCTGGTGGTGGT  
CTGGTGGCGGCTCTGAGGGTGGTGGCTCTGAGGGTGGCGGTTCTGAGGGTGGCGGCTCTGAGGGAGGCGGT **CATAGTAC** CCGGTGGTGGC  
TCTGGTCCGGTGATTTGATTATGAAAAGATGGCAAACGCTAATAAGGGGGCTATGACCGAAAATGCCGATGAAAACGCGCTACAGTCTGACGCT  
AAAGGCAAACCTGATTCTGCTGCTACTGATTACGGTGTCTGCTATCGATGGTTTCATTGGTGACGTTCCGGCCTTGCTAATGGTAATGGTGTACTGGTG  
ATTTGCTGGCTCTAATCCCAAATGGCTCAAGTCCGTGACGGTGATAATCA **ACAAAATAAAG** AATTCCGTCAATATTACCTCCCTCCCTCAATCGG  
TTGAATGTCGCCCTTTTGTCTTTGGCGCTGGTAAACCA **ATGAATTTCTA** TTGATTGTGACAAAATAAACTTATCCGTGGTGTCTTTGCGTTCTTTTATATGTTG  
CCACCTTATGTATGATTTTCTACGTTTGTAACTACTGCGTAATAAGGAGTCTAATCATGCCAGTCTTTTGGGTATTCCGTTATTATTGCGTTTCTCCTCGGT  
TCCTTCTGGTAACTTTGTTCCGGT **TTAATAAAA** ATCTGCTTACTTTTCTTAAAAAGGGCTTCGGTAAGATAGCTATTGCTATTTCAITGTTCTTGTCTTATTATTG  
GGCTTAACTCAATCTTGTGGGTATCTCTCTGATATTAGCGCTCAATTACCCTCTGACTTTGTTACAGGGTGTTCAGTTAATTCTCCCGTCTAATGCGCTTCCC  
TGTTTTATGTTATTCTCTCTGTAAGGCTGCTATTTTCATTTTGACG **AGAGGG** TTAACAATAAAATCGTTTCTATTGGATTGGGATAATAATATGGC **TGTT**  
**ATTTGT** AACTGGCAAATTAGGCTCTGGAAAGACGCTCGTTAGCGTTGGTAAGATTCAGGATAAAATTGTAGCTGGGTGCAAATAGCAACTAATCTTG  
ATTAAGGCTCAAACCTCCCGCAAGTCGGGAGGTTGCTAAAACGCCTCGCGTCTTAGAATACCGGATAAGCCCTCTATATCTGATTTGCTTGCTA  
TTGGGCGCGGTAATGATTCCTACGATGAAAATAAAAACGGCTTGTGTTCTCGATGAGTGCGGTACTTGGTTAATACCCGTTCTTGGAAATGATAAGG  
AAAGACAGCCGATTATTGATTGGTTTCTACATGCTCGTAAATTAGGATGGGATATTATTTTCTGTTTCAAGGACTTATCTATTGTTGATAACAGGCGCGTCT  
GCATTAGCTGAACATGTTGTTATTGTCGTCGCTCTGGACAGAATTACTTTACCTTTTGTCCGTACTTATATTCTTATTACTGGCTCGAAAATGCCTCTGCCT  
AAATTACATGTTGGCGTTGTTAAATATGGCGATTCTCAATAAGCCCTACTGTTGAGCGTTGGCTTATACTGGTAAGAATTTGATAACGCATATGATACTA  
AACAGGCTTTTTCTAGTAATTATGATTCCGGTGTATTCTTATTAACGCCTTATTATCACACGGTCCGGTATTCAAACCATAAATTTAGGTCAGAAGATGA

Specific interactions: does the transition change?





# Unresolved questions

Does linear sDNA fold?

How does local/cooperative is the process?

Is there a DNA theta point?

How important is topology?

Thanks for the attention!

*And they lived happily ever after*