

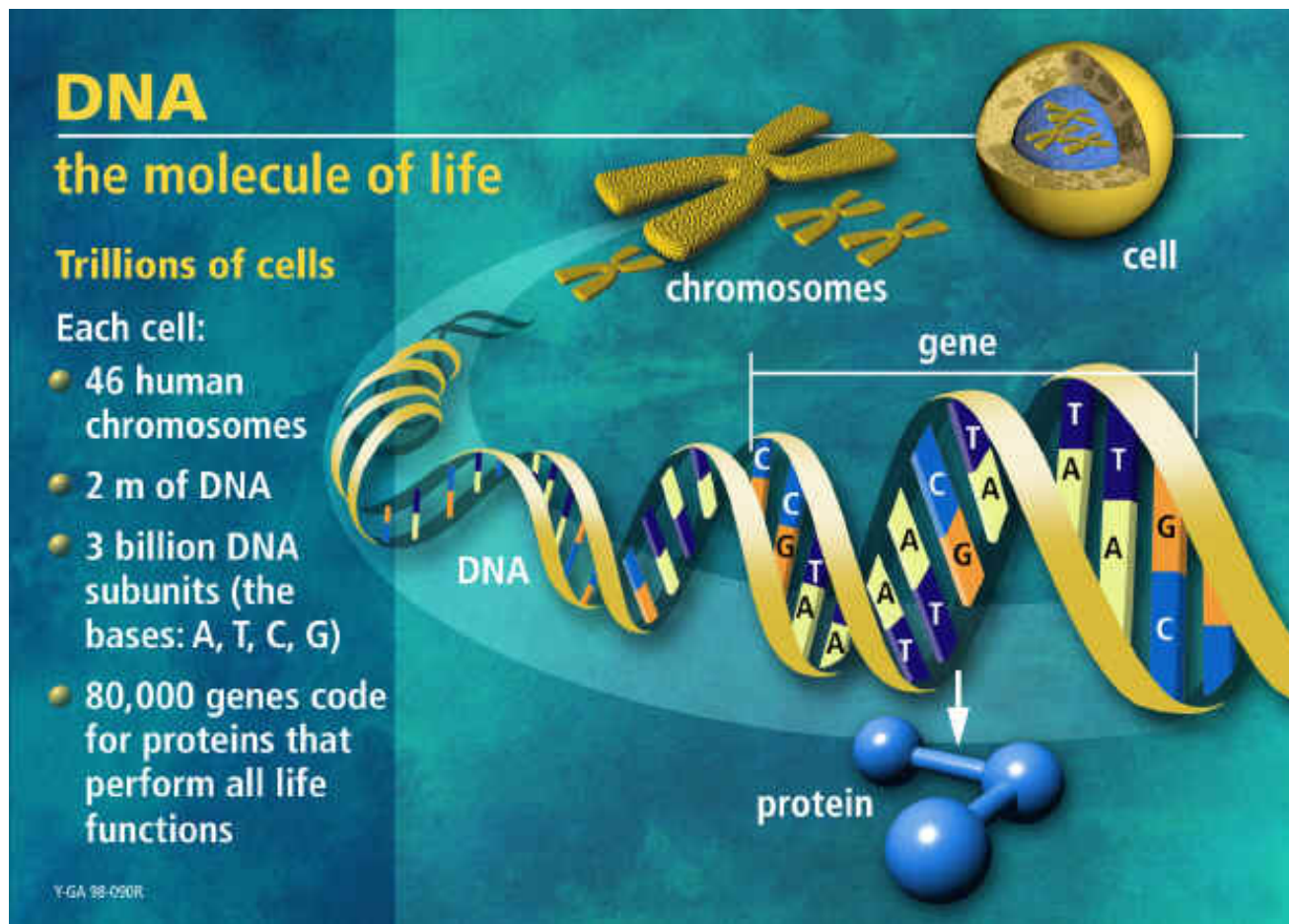
Multiscale Modeling of Structural and Electronic Properties of DNA

- Motivation: sequencing, epigenetics, replication and repair, ...
- Coarse-grained potential: model, simulations, validation
- Translocation dynamics through nanopores
- Electronic properties

EMRS - May 14-17, 2012, Strasbourg

Symposium J: DNA Directed Programmable Self-Assembly of Nanoparticles into Meta Materials for Energy and other Applications



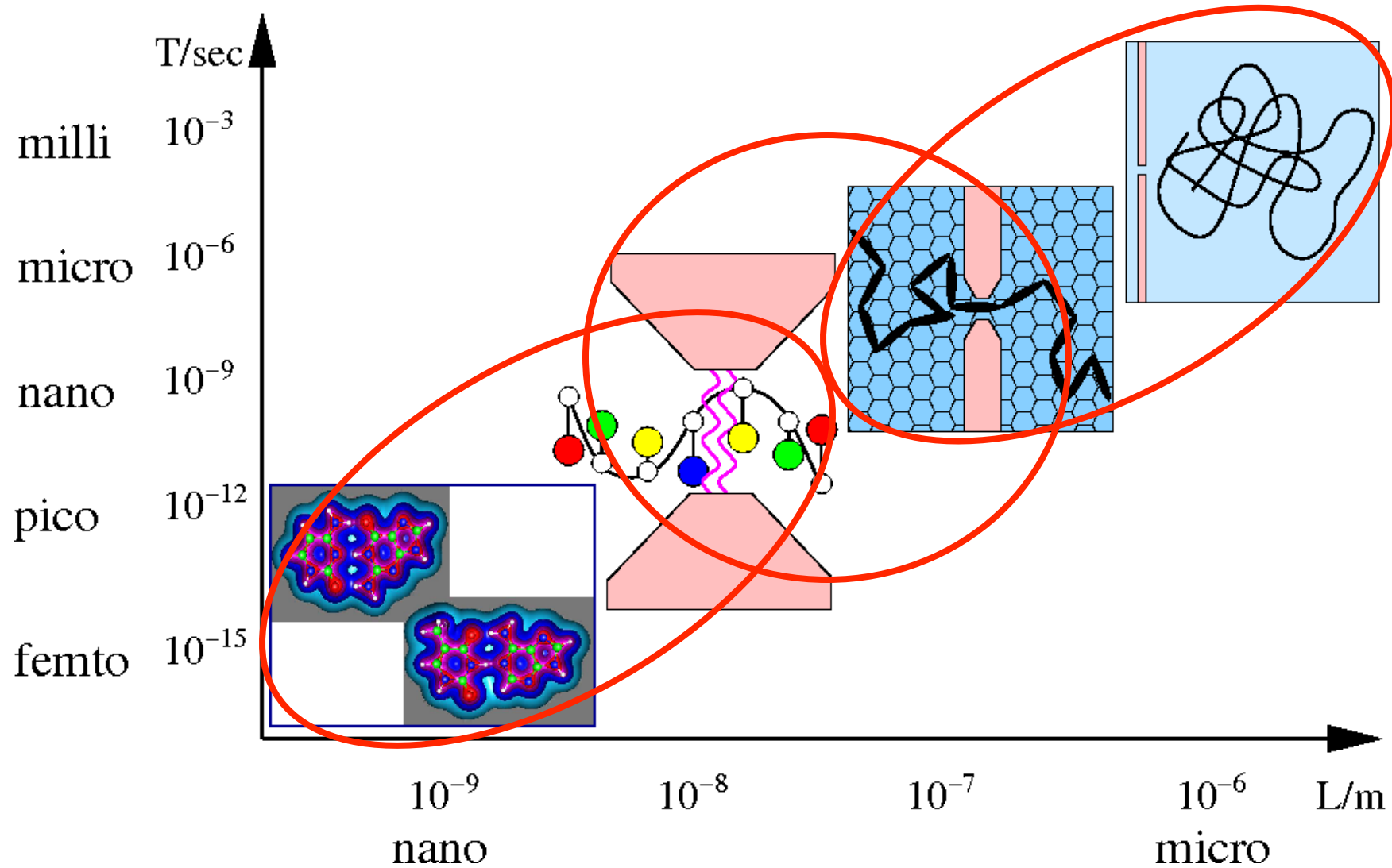


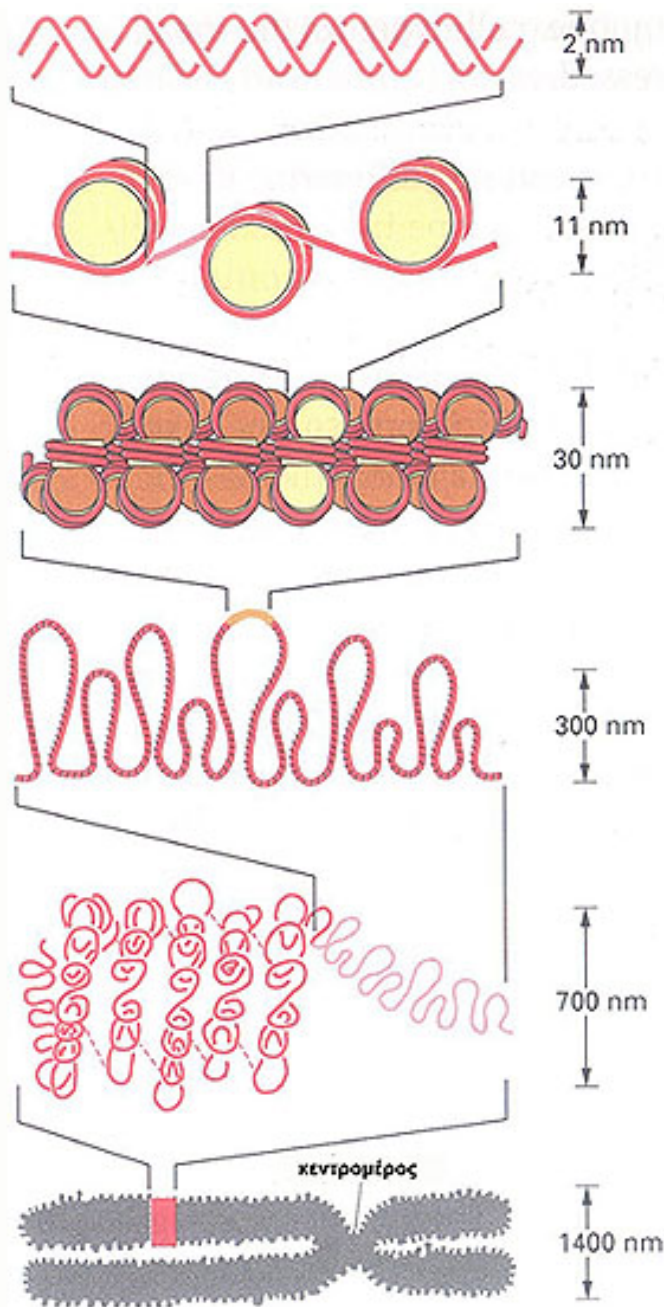
DNA sequencing: biochemical methods - cut pieces (ending at specific bases), measure by **gel electrophoresis**.

Instead, use **electronic signature** for sequencing - multiscale process

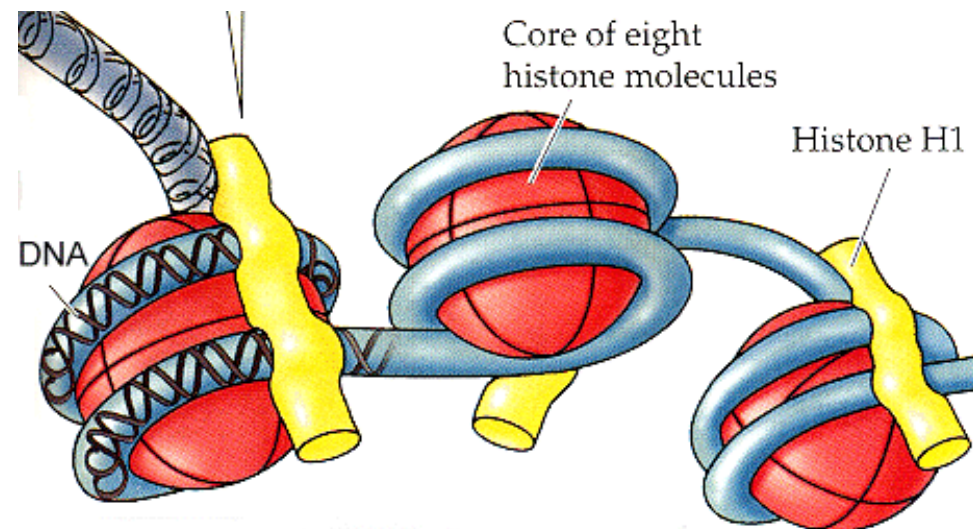


Multiple length/time scales: DNA electronic sequencing





Structure of DNA on several scales: From NUCLEOSOME to CHROMOSOME

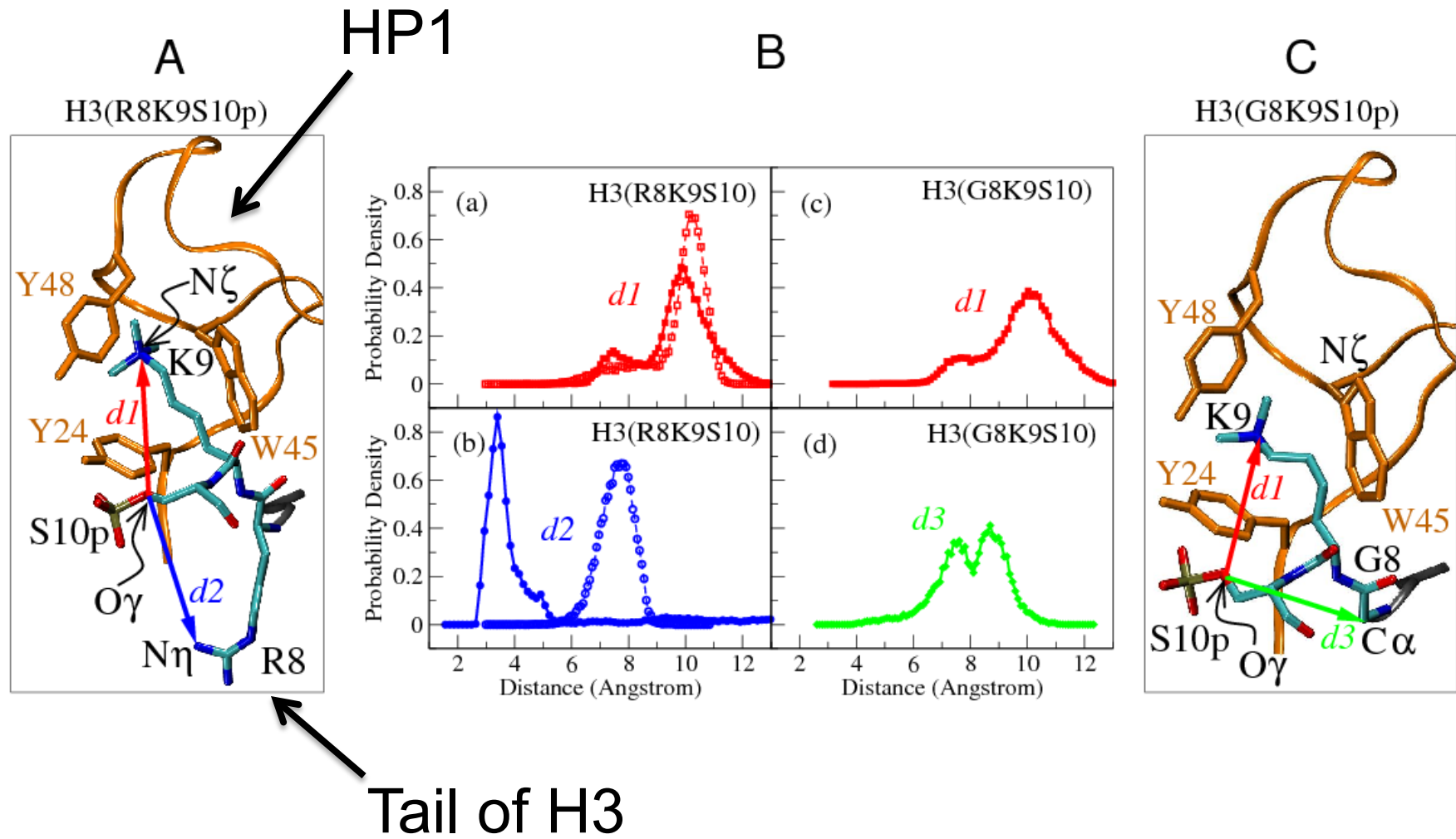


EPIGENETICS: passing genetic information **NOT** encoded in DNA base sequence



Papamokos *et al.*,
Biophysical J.
(April 2012)



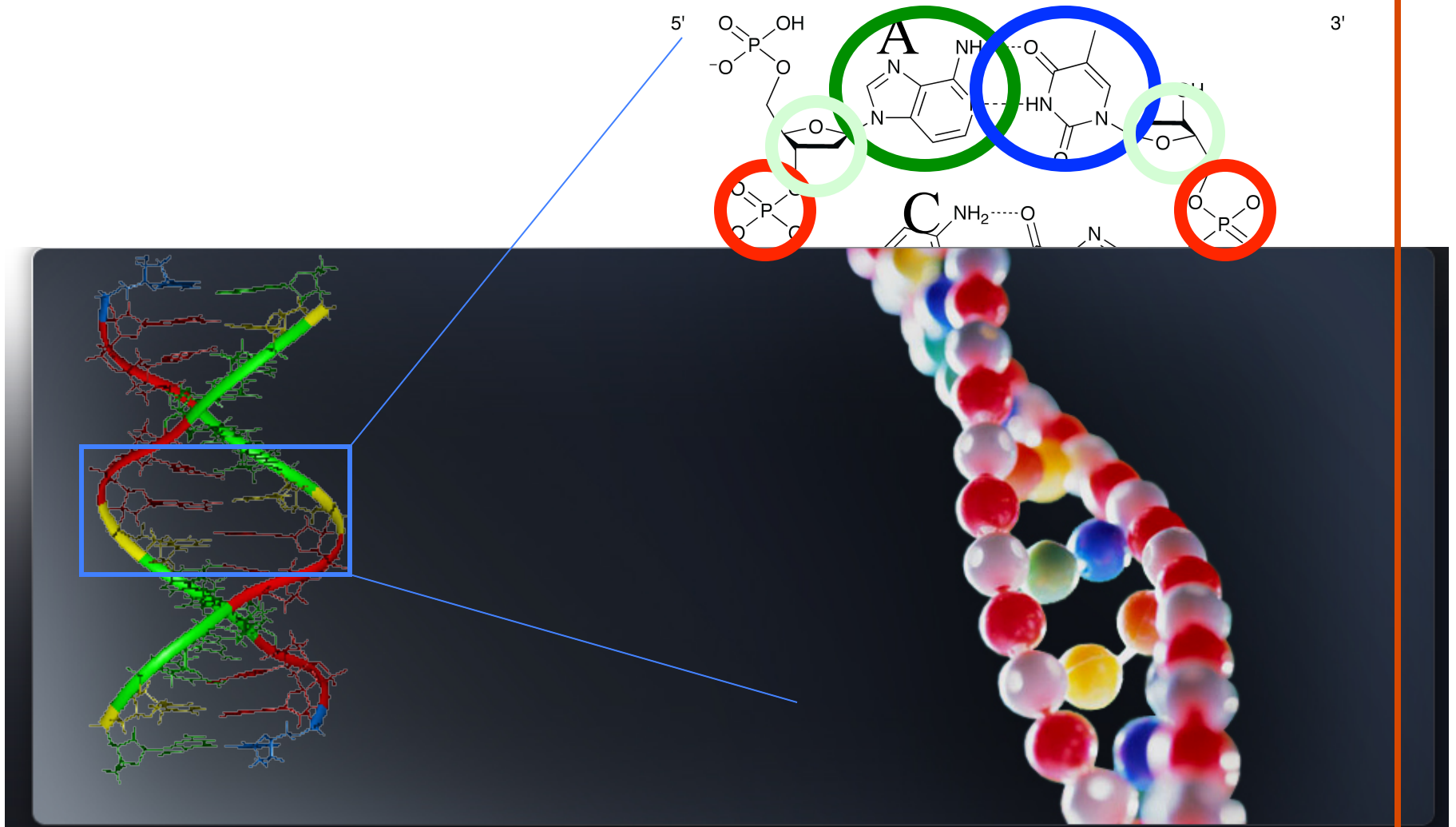


Papamokos *et al.*, Biophysical J. (2012)



Coarse-grained potential fo DNA

Deoxyribonucleic Acid



W. Hsu, M. Fyta, G. Lakatos, S. Melchionna, EK (2012)



Goal: derive coarse-grained potential:
minimal (but sufficient) model;
all parameters from *ab-initio* calculations*.

Assumption: **separable** interactions

- Hydrogen bonding (distance; dihedral, flip angles)
- Stacking interactions (distance; twist angle)
- Backbone interactions (distance; 3'-5' orientation)
- Electrostatic interactions

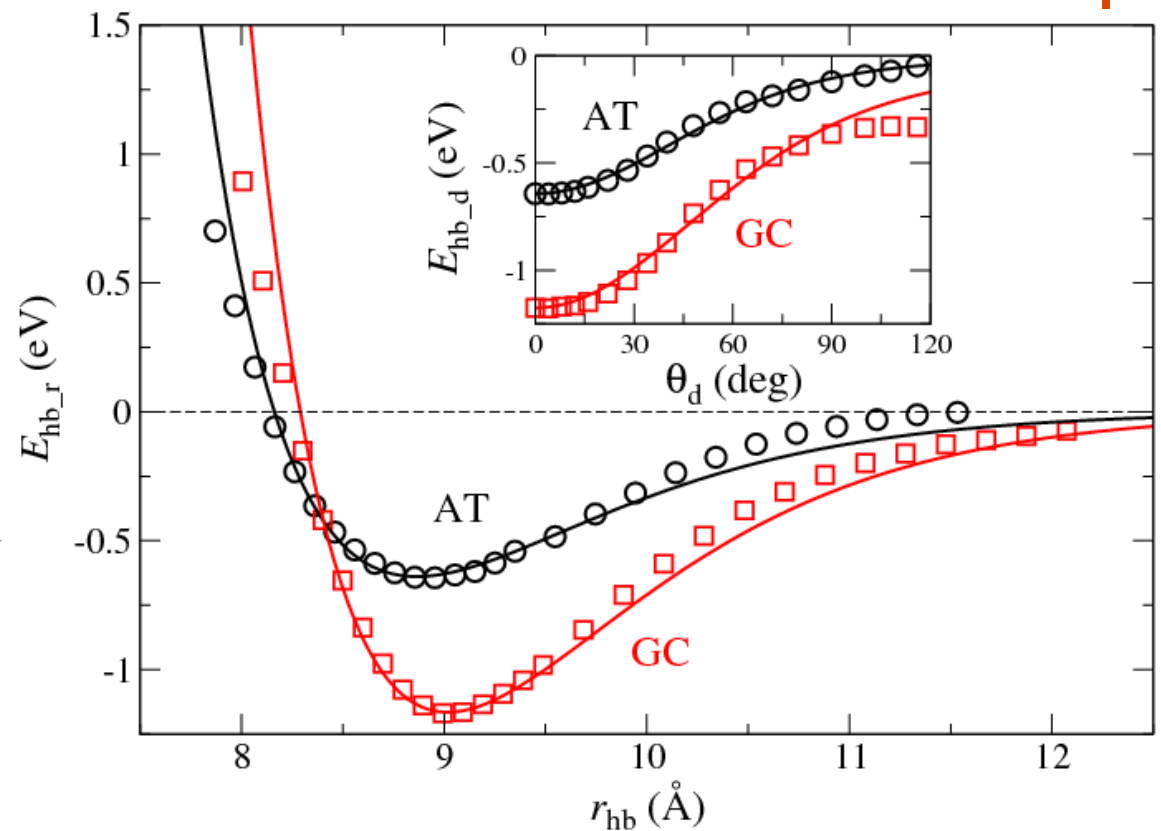
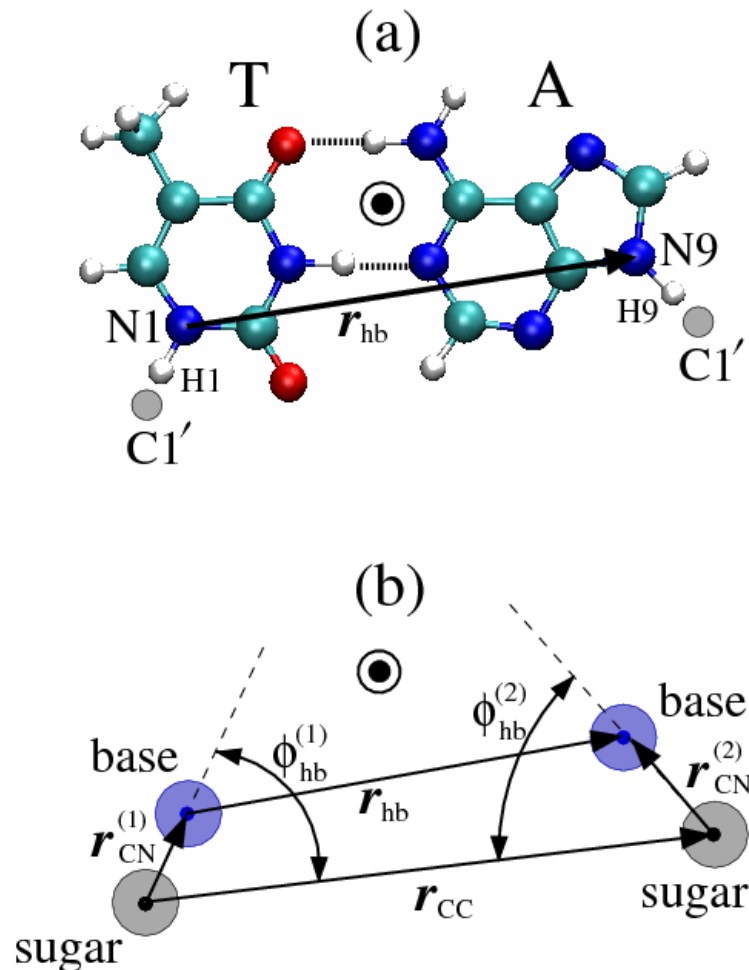
* To the extent possible

Data points: DFT calculations,
Lines: fits with simple curves

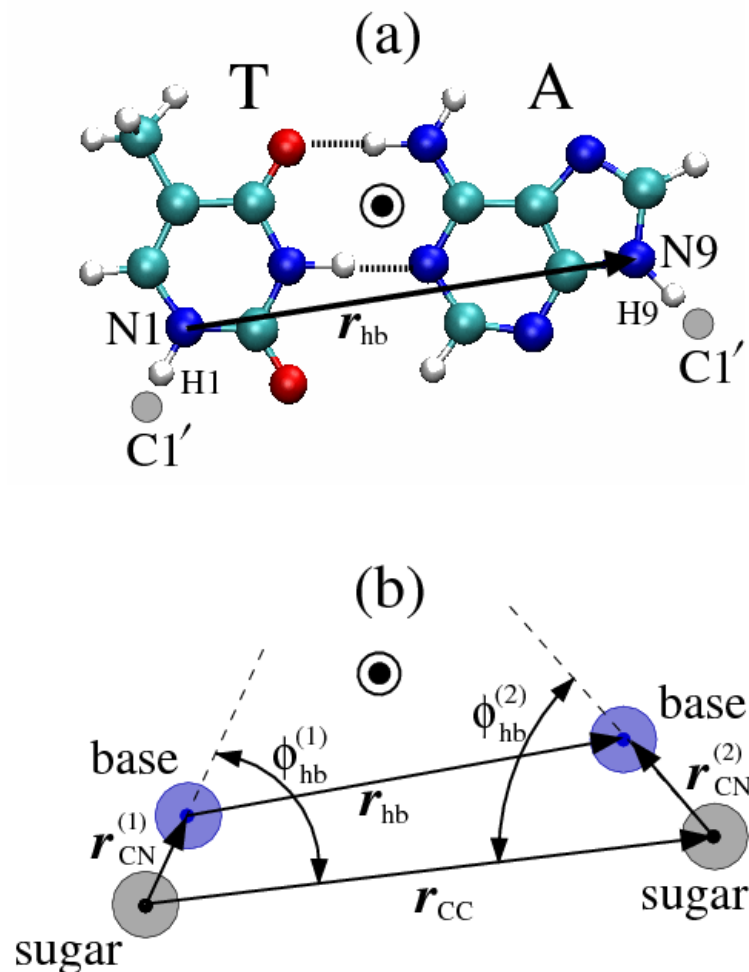


Coarse-grained potential – hydrogen bonding I

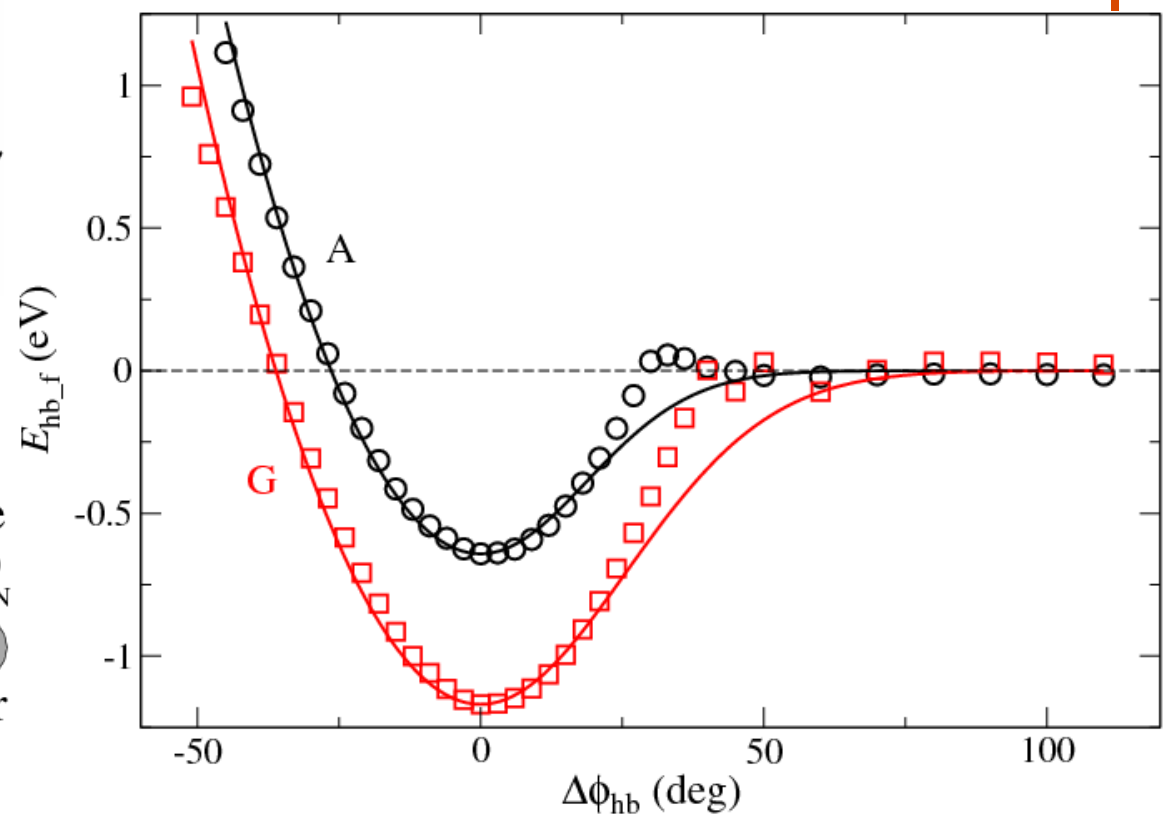
Distance between bases,
dihedral angle



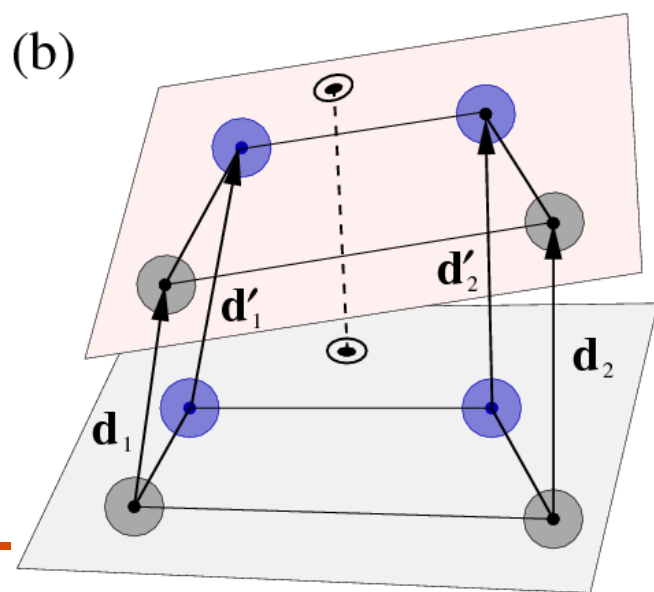
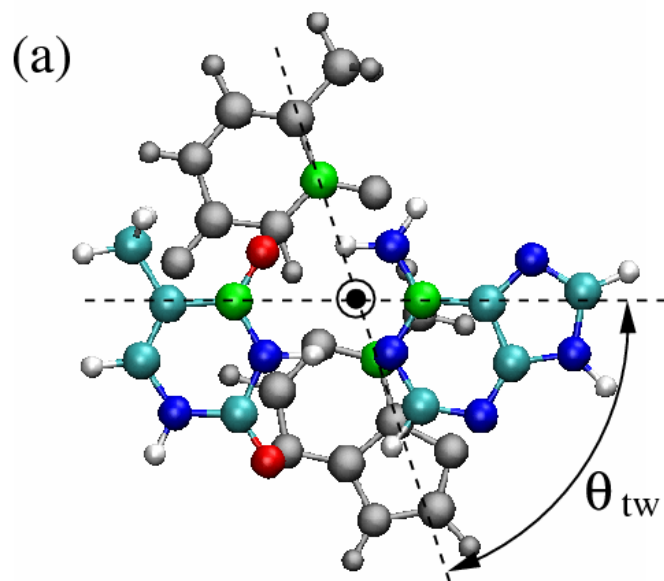
Coarse-grained potential – hydrogen bonding II



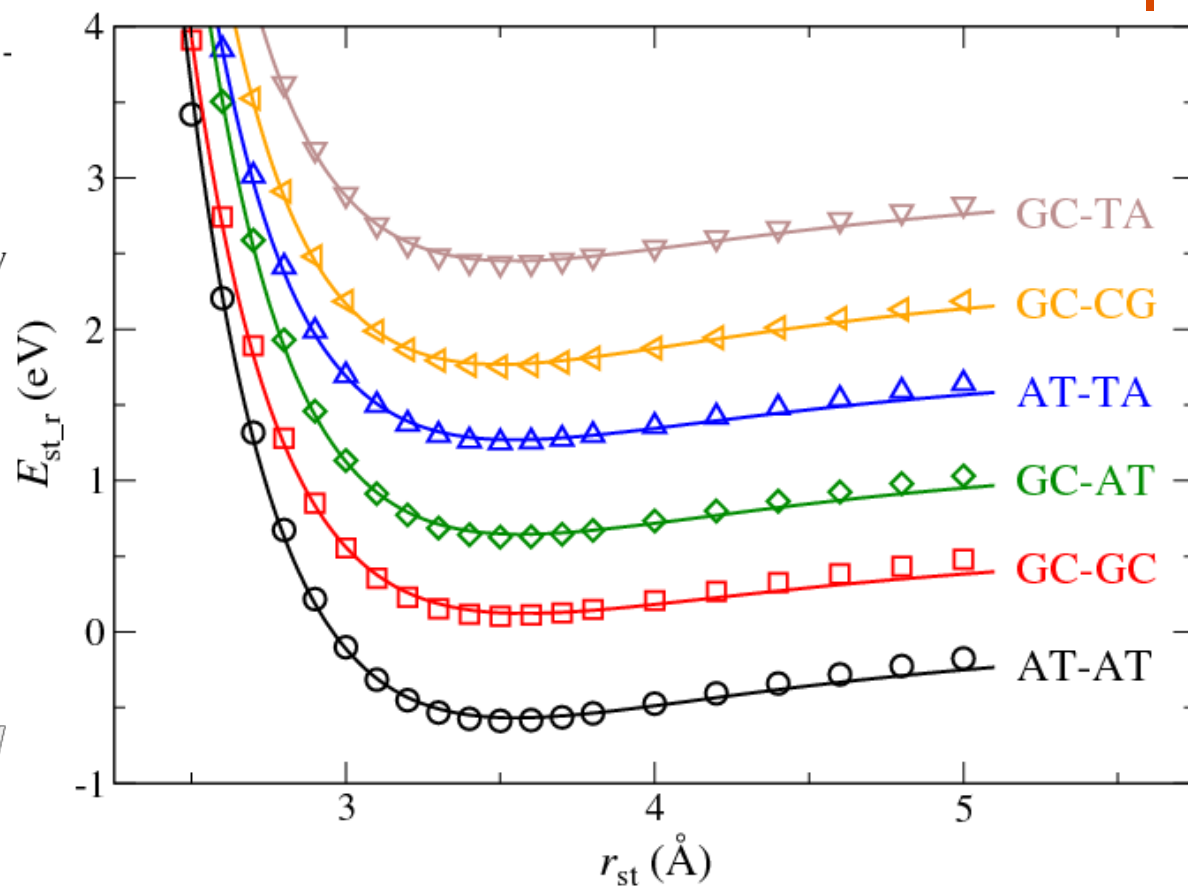
Flip angle



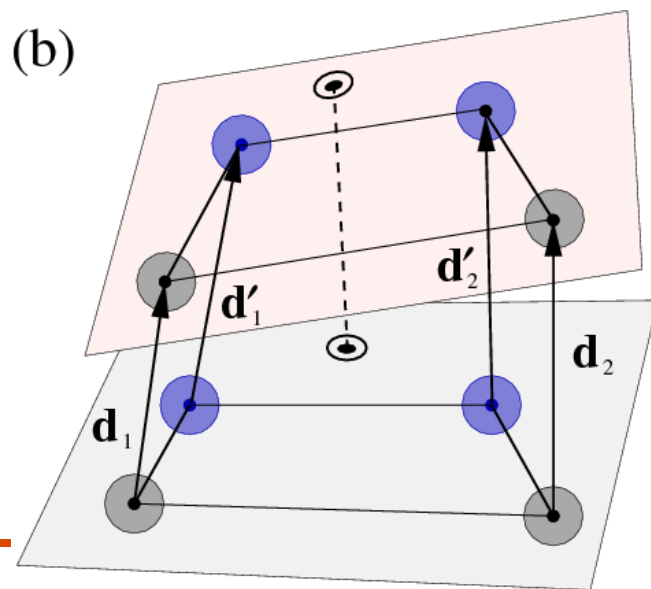
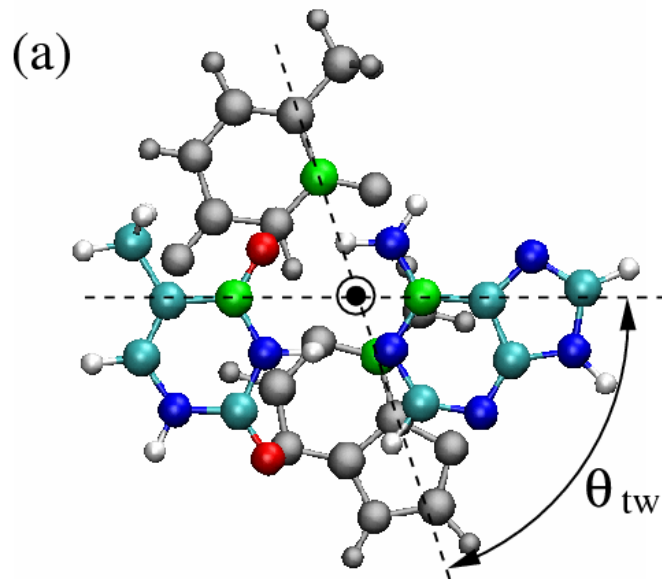
Coarse-grained potential – stacking interactions



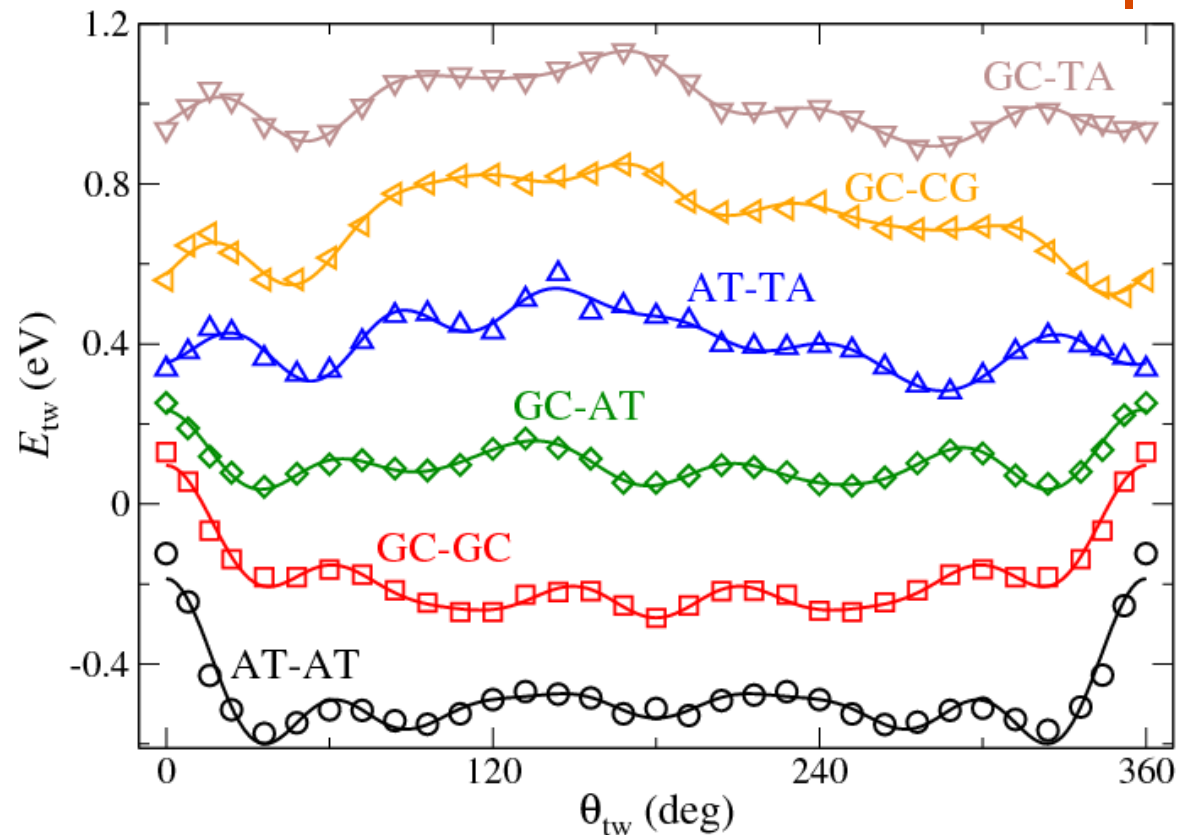
Distance between planes



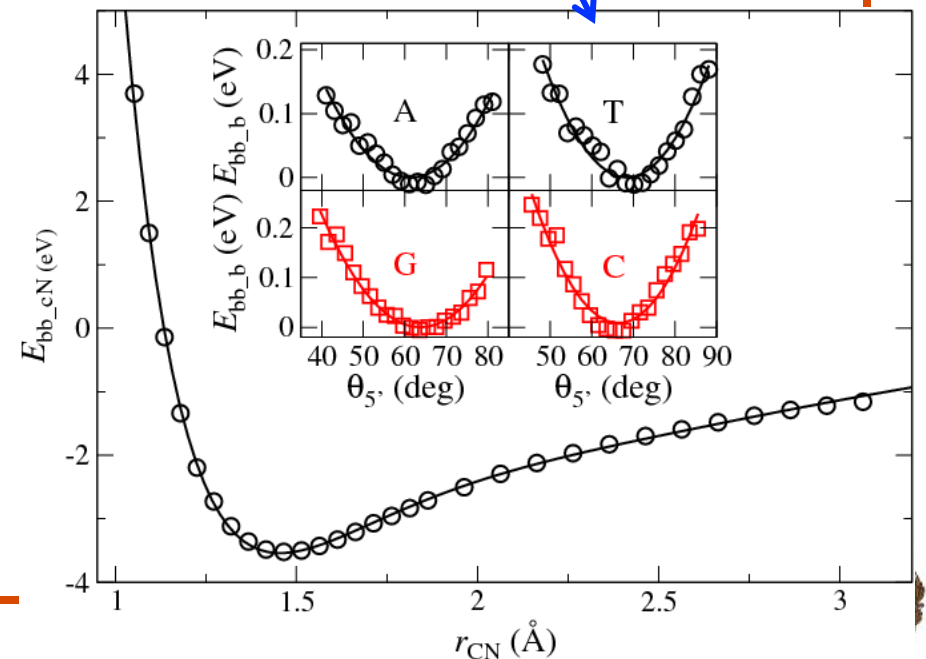
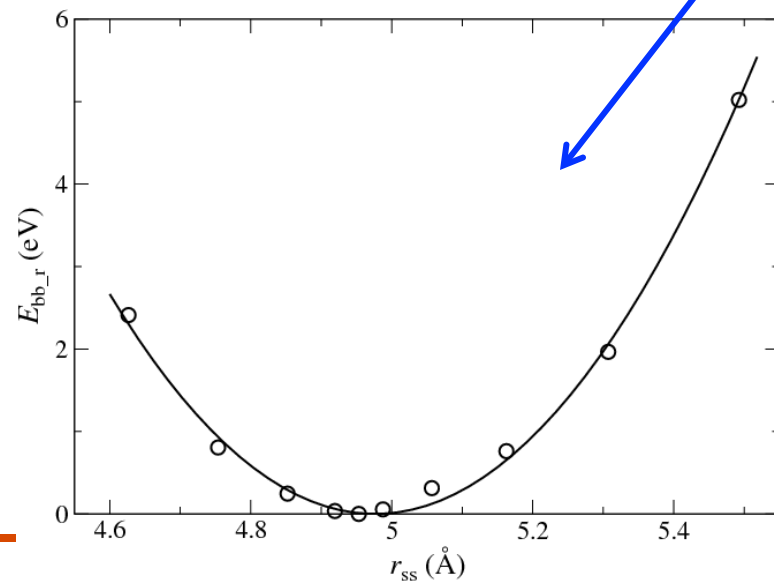
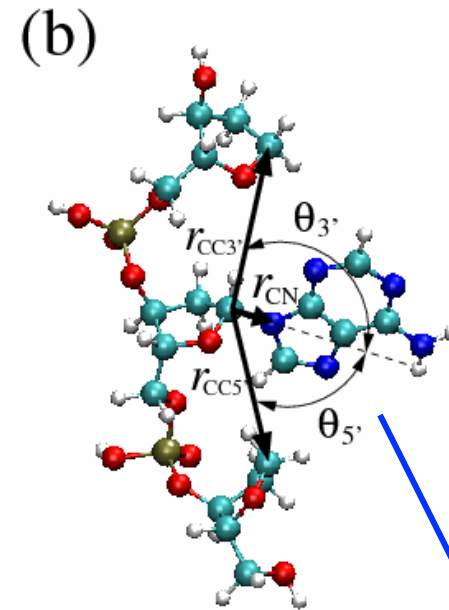
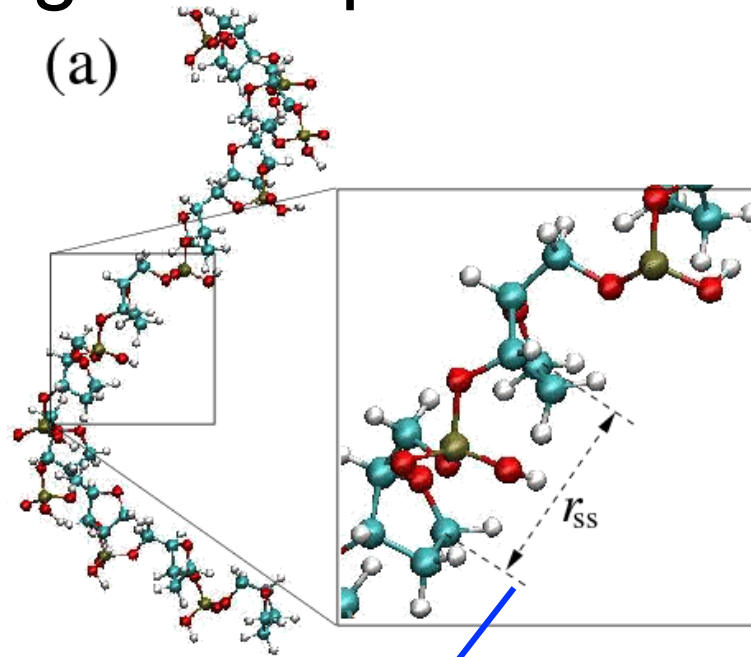
Coarse-grained potential – stacking interactions



Twist angle between pairs

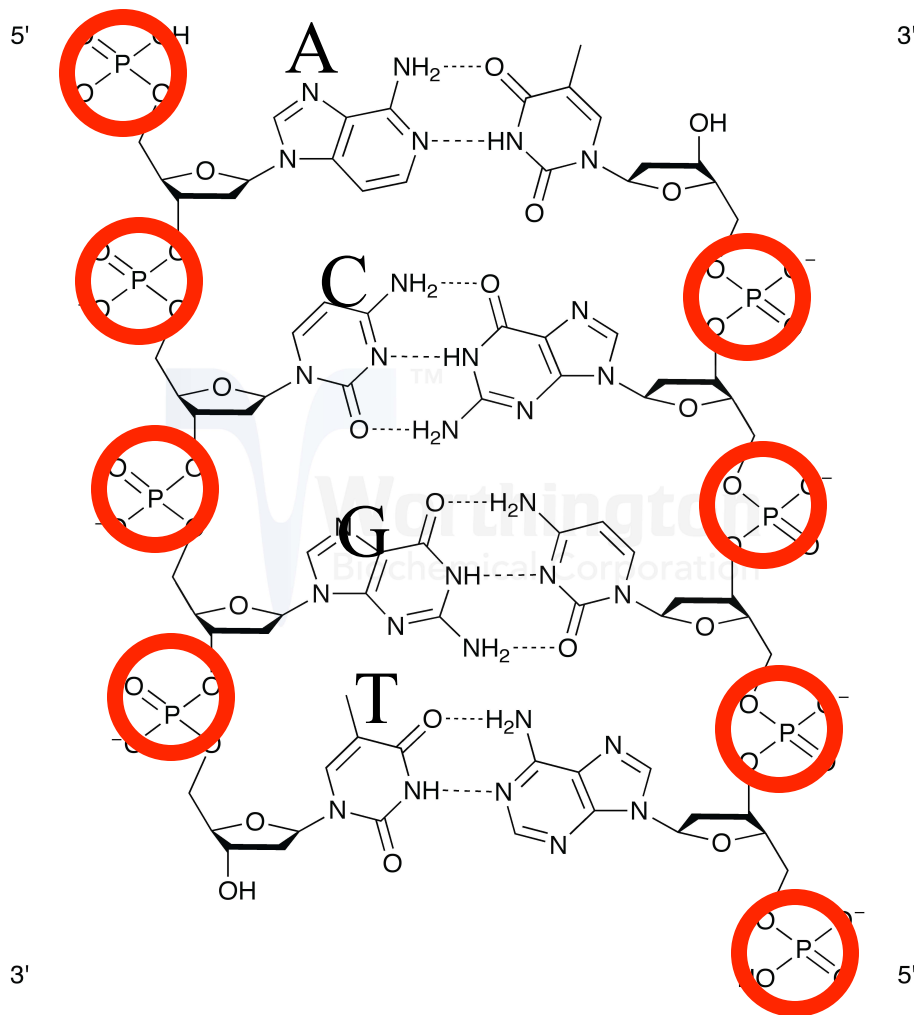


Coarse-grained potential – backbone interactions



Coarse-grained potential – electrostatic interactions

Deoxyribonucleic Acid



5'-Adenine-Cytosine-Guanine-Thymine-3'

$$E = \frac{1}{4\pi\epsilon_0\epsilon(r)} \frac{e^2}{r}$$

$$\epsilon(r) = \epsilon_{\text{in}} \quad r < r_0$$

$$\epsilon(r) = \epsilon_{\text{in}} e^{\alpha(r-r_0)} \quad r_0 < r < r_1$$

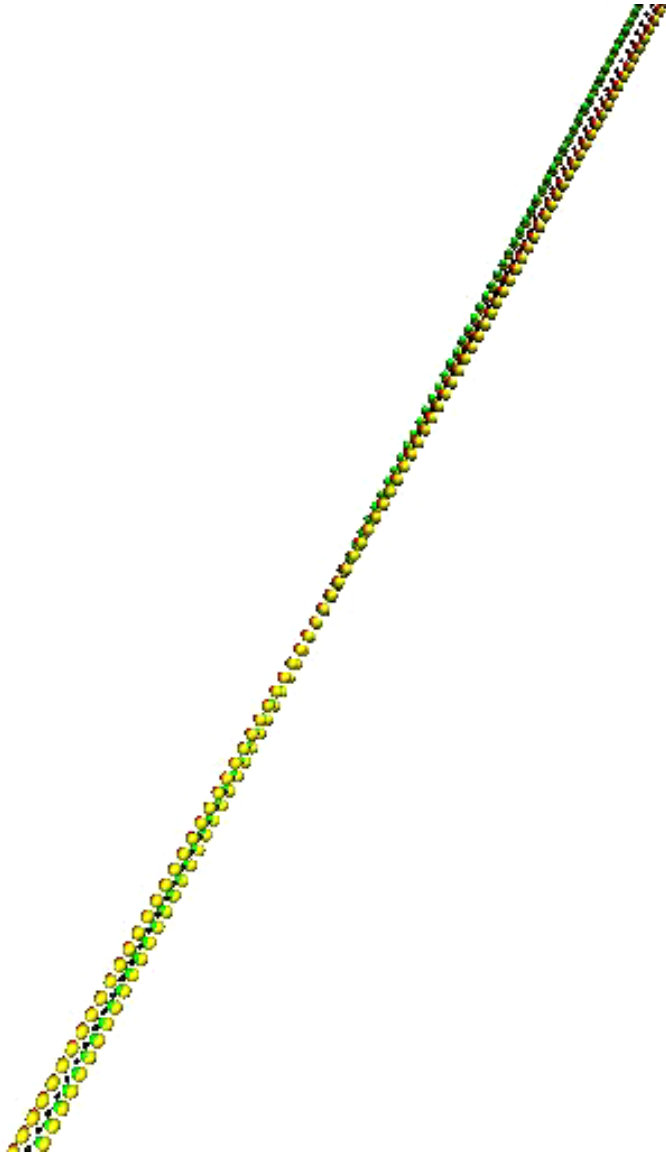
$$\epsilon(r) = \epsilon_{\infty} e^{-\kappa r} \quad r > r_1$$

$$\kappa^{-1} = \sqrt{\frac{\epsilon_0 \epsilon_{\infty} k_B T}{2 N_A e^2 I}}$$

Three parameters !



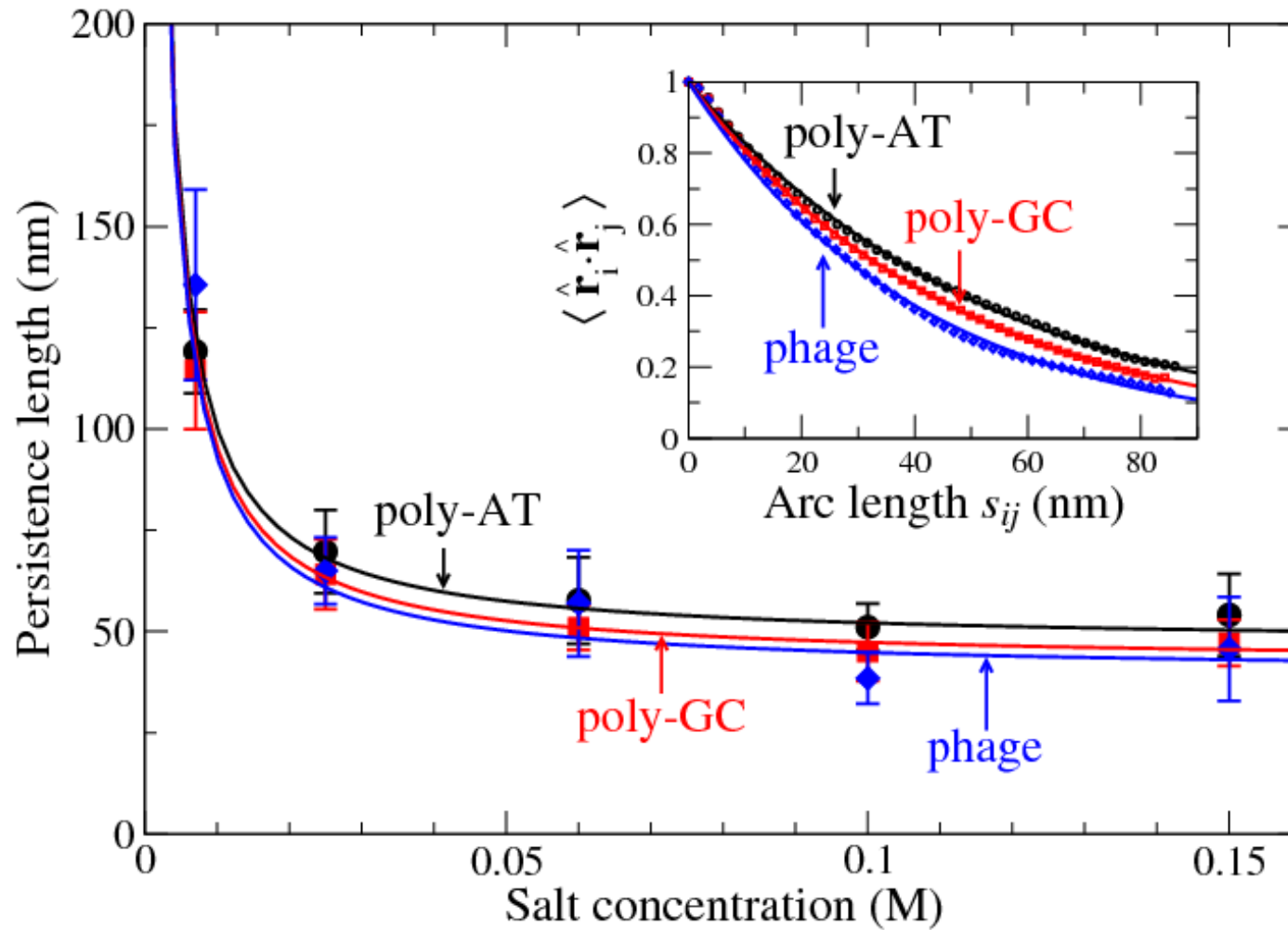
Coarse-grained potential – validation



Two parallel strands
coil to form double-helix



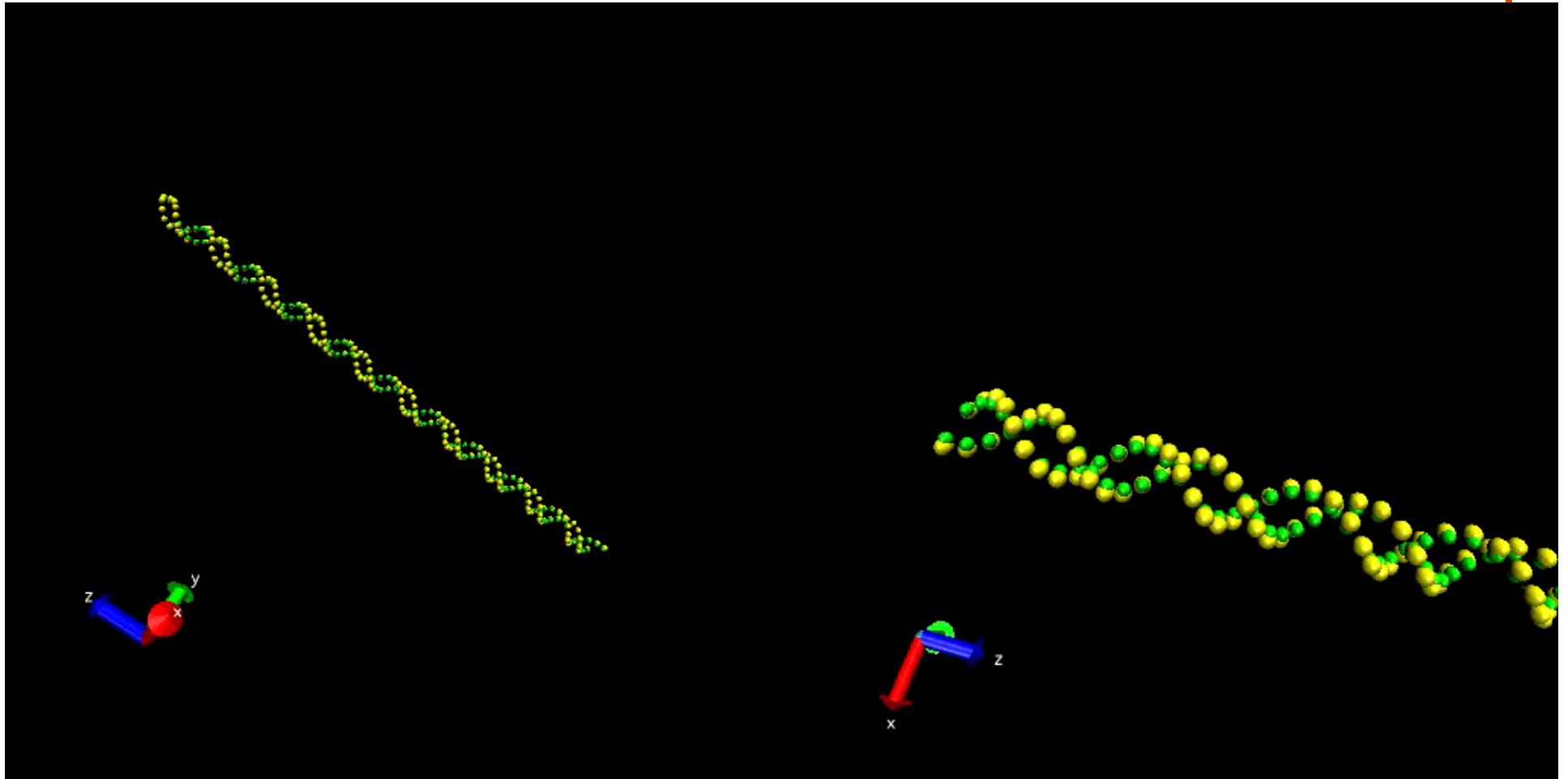
Coarse-grained potential – validation



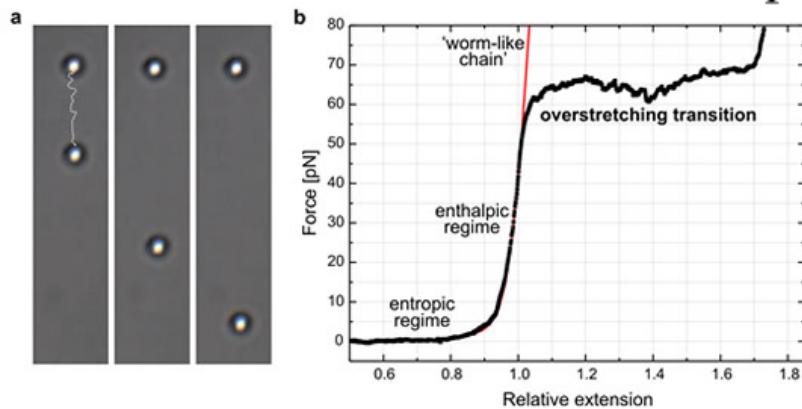
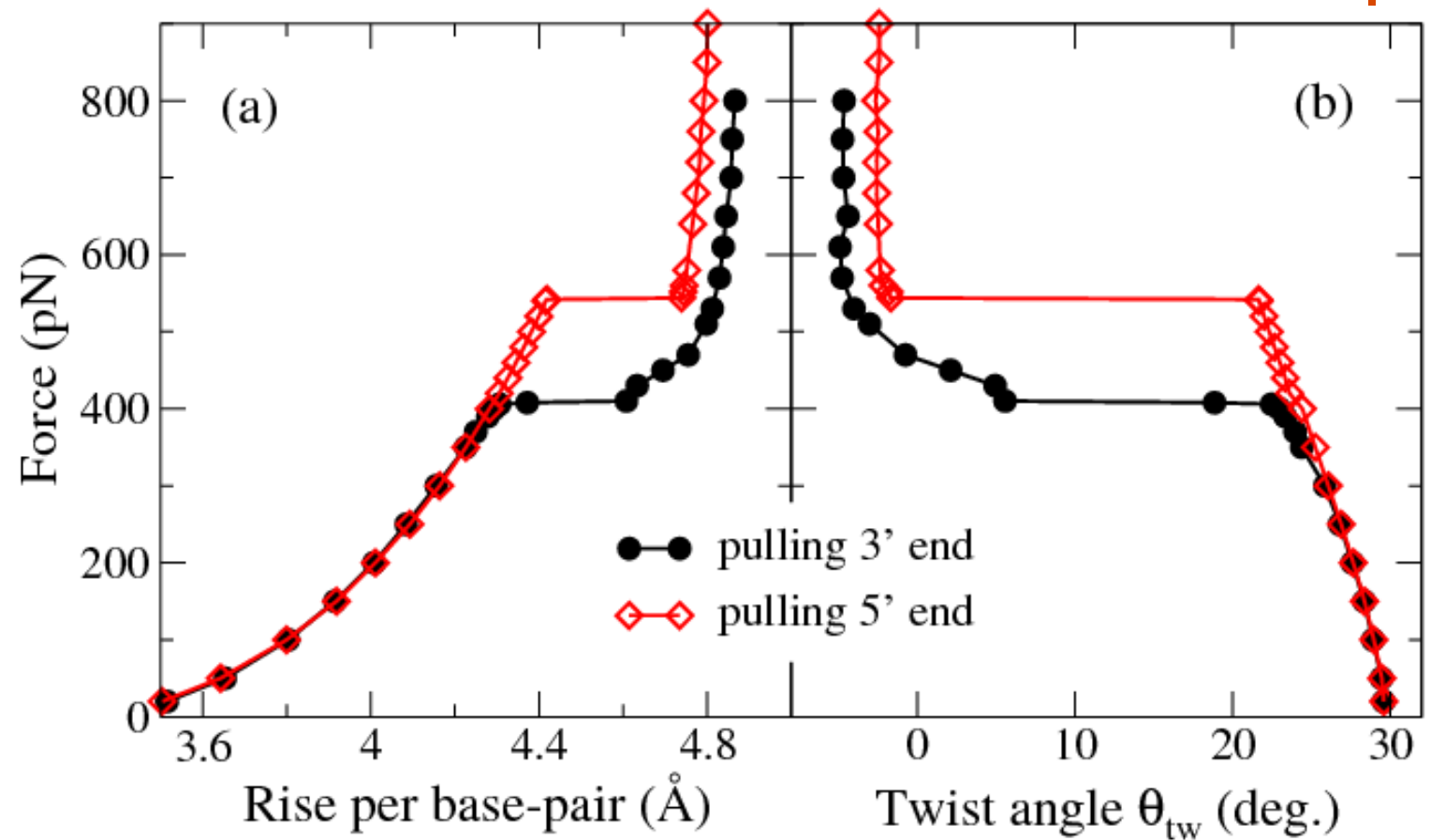
Persistence length (~50 nm)



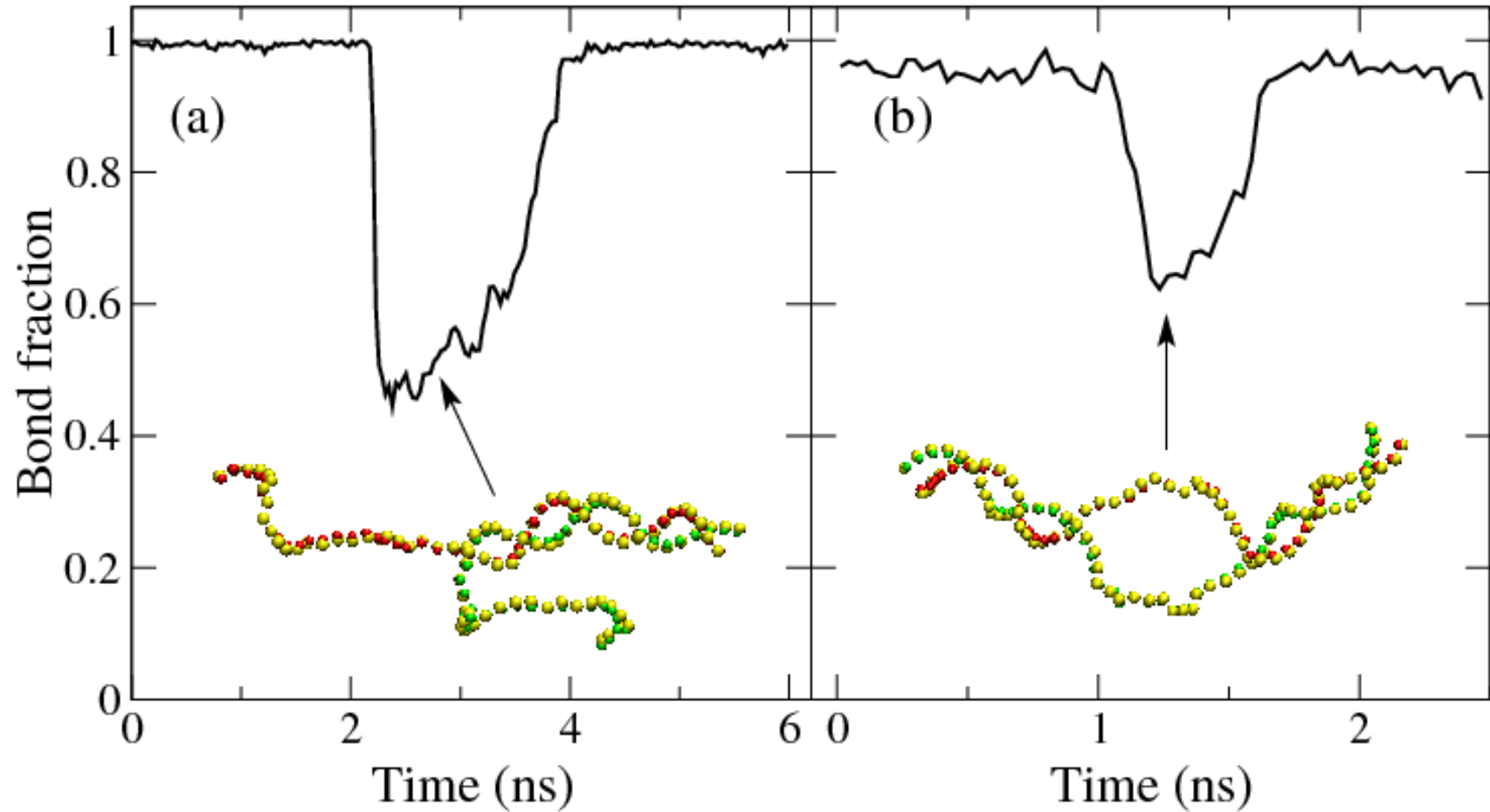
Force-extension simulations



Force-extension simulations



Melting simulations



Students: Chia Wei Hsu
Ryan Barnett
Ari Turner

Postdocs: Maria Fyta
Greg Lakatos
Simone Mechlionna
Paul Maragakis
Gang Lu
Sheng Meng

Visitors: Sauro Succi
Costas Papaloukas
George Papamokos

