

Biomolecular Simulation

BT2123

Lecture 5 : Creating an all-atom model of DNA

Himanshu Joshi 30 January 2024



- Historical perspective
- Foundations of Molecular Mechanics (MM)
- **Statistical ensembles**
- Quantum Mechanics (QM)
- Introduction molecular dynamics simulations
- Equation of motion,
- Force-fields, Scheme of integrations,
- Langevin Dynamics,
- Non-bonded Computations,
- Brownian Dynamics,
- Monte Carlo Techniques,
- Coarse Graining Models

Reading material

Who discovered DNA ?

<https://www.nature.com/scitable/topicpage/discovery-of-dna-structure-and-function-watson-397/>

Friedrich Miescher, 1869

doi:10.1006/jmbi.2001.4987 available online at <http://www.idealibrary.com> on IDEAL® *J. Mol. Biol.* (2001) **313**, 229–237

JMB



NOMENCLATURE

A Standard Reference Frame for the Description of Nucleic Acid Base-pair Geometry

**Wilma K. Olson, Manju Bansal, Stephen K. Burley
Richard E. Dickerson, Mark Gerstein, Stephen C. Harvey
Udo Heinemann, Xiang-Jun Lu, Stephen Neidle, Zippora Shakked
Heinz Sklenar, Masashi Suzuki, Chang-Shung Tung, Eric Westhof
Cynthia Wolberger and Helen M. Berman**

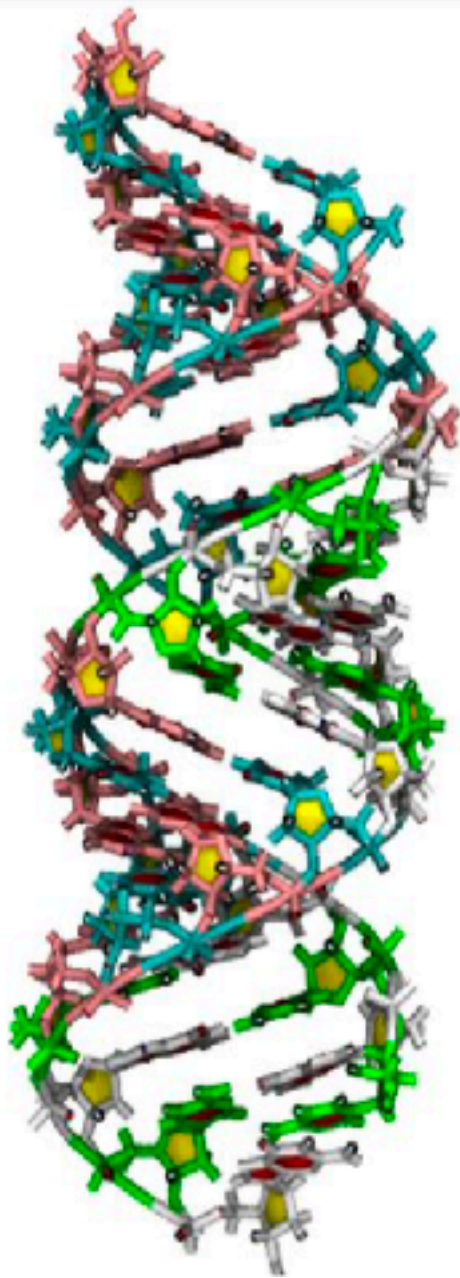
© 2001 Academic Press

Keywords: nucleic acid conformation; base-pair geometry; standard reference frame

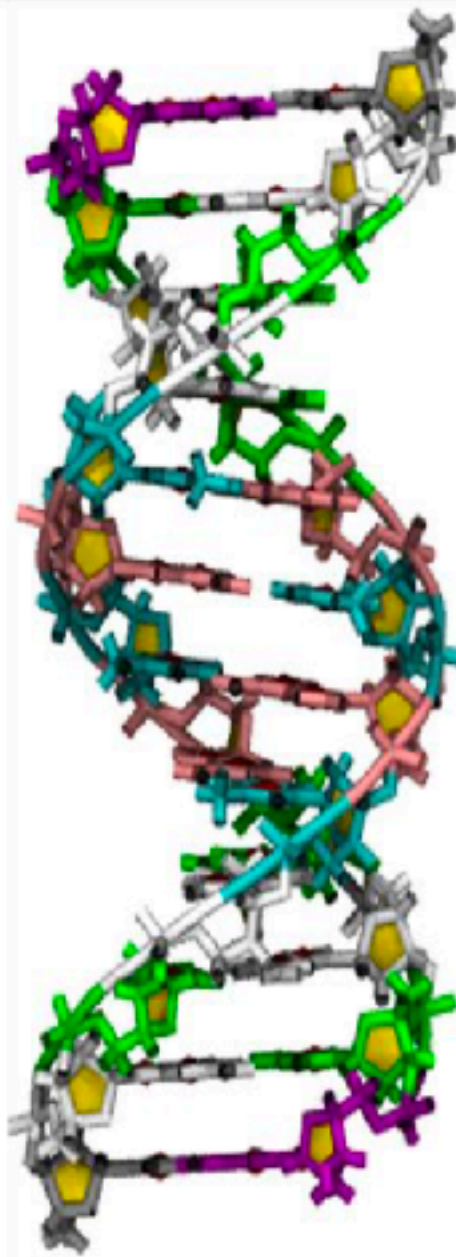
Various forms of DNA



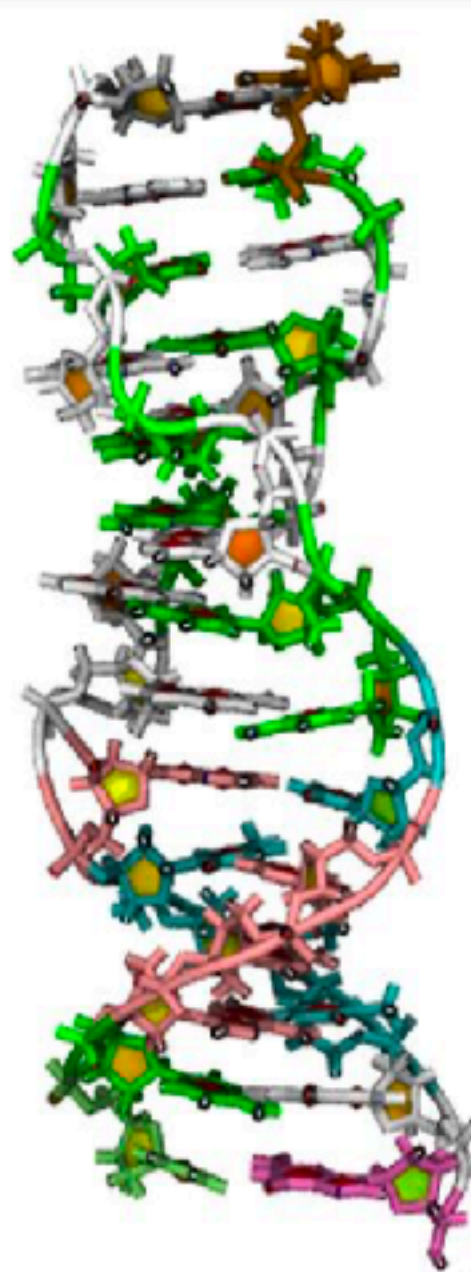
ssDNA



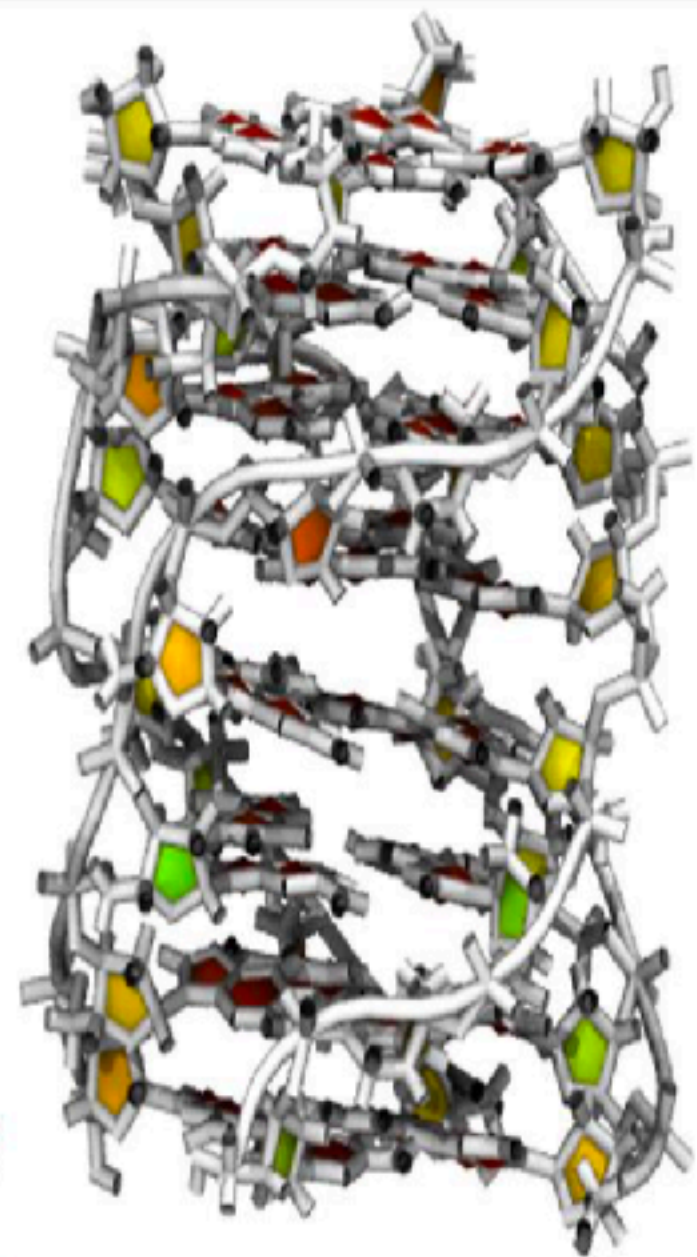
A-DNA



B-DNA
(canonical)



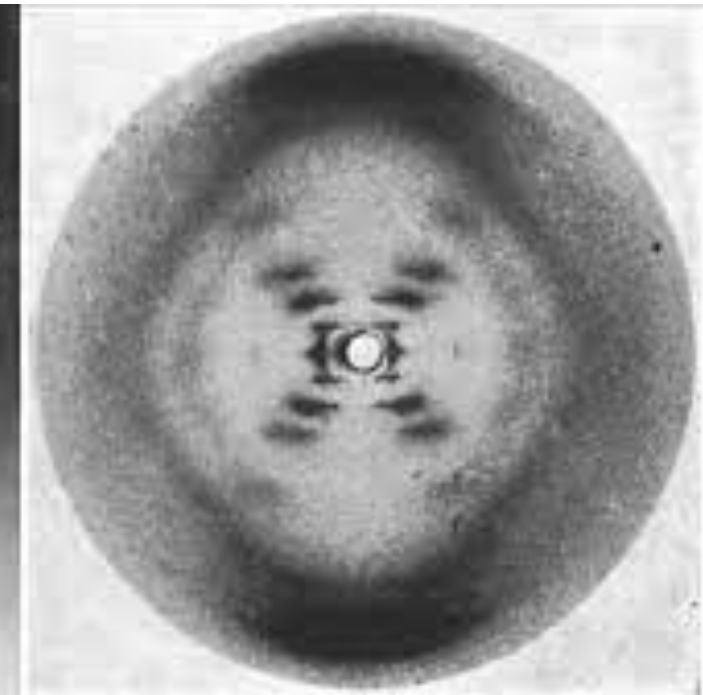
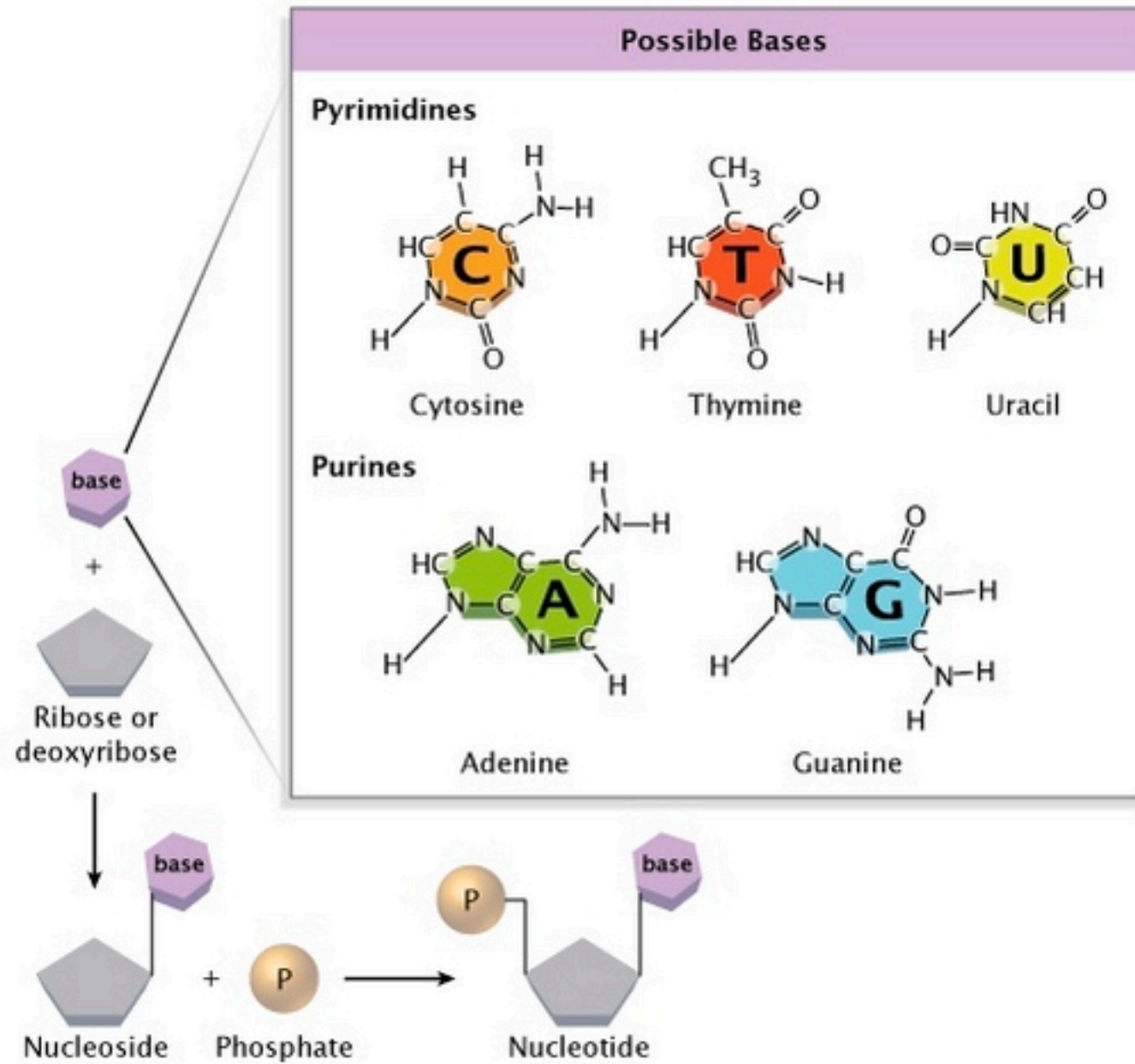
Z-DNA



G-quadruplex

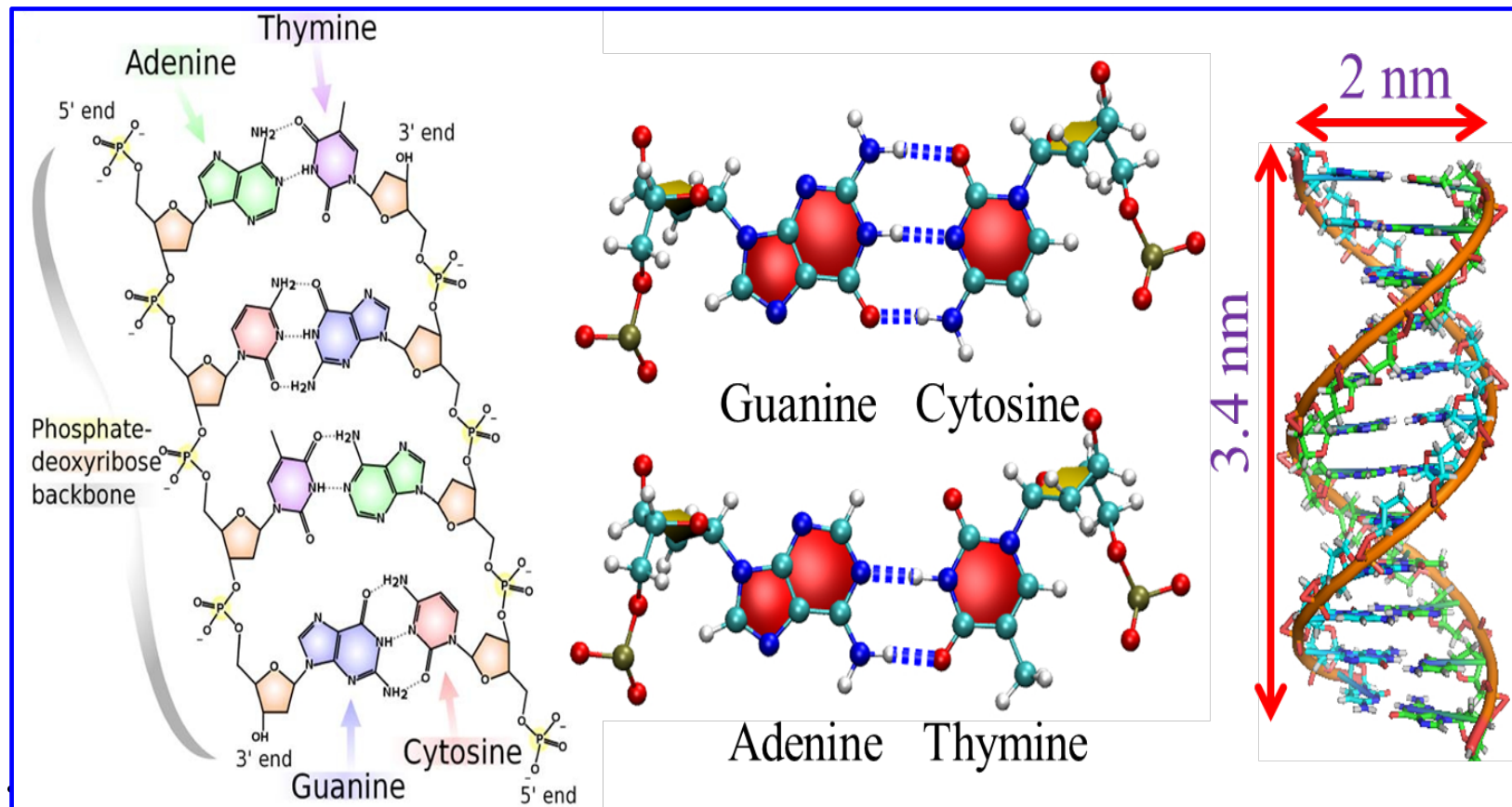


3 D structure of DNA



Structural DNA Nanotechnology

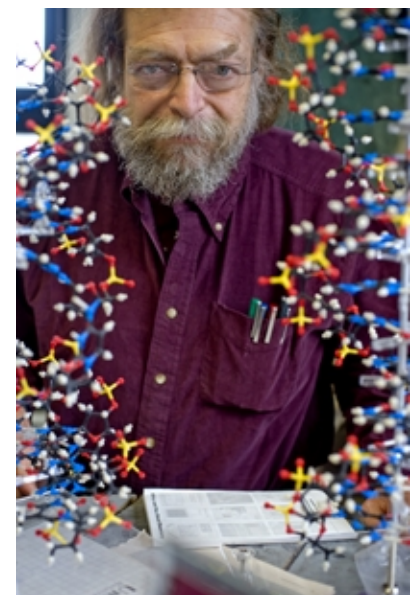
- There is plenty of room at the bottom.
- Bottom up self-assembly : DNA as a construction material



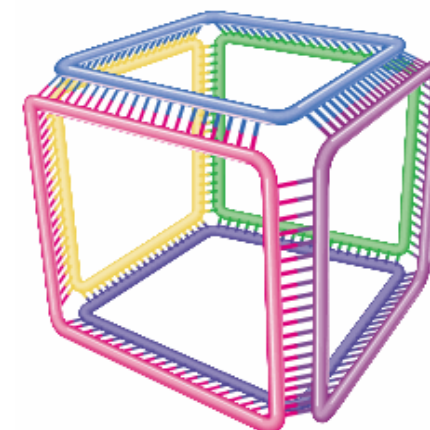
Why DNA

1. Molecular recognition property
2. Chemically easy to modify the backbone
3. Well studied molecule.

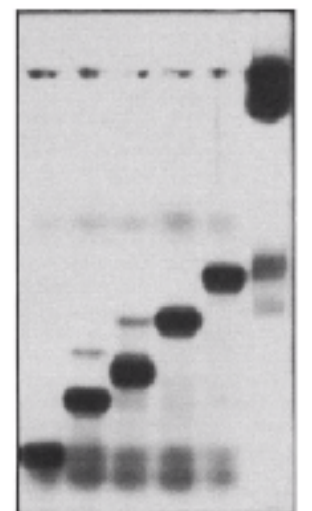
R. Feynman




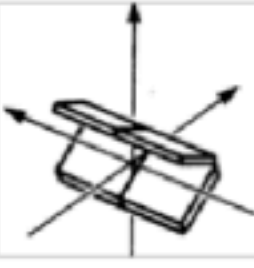
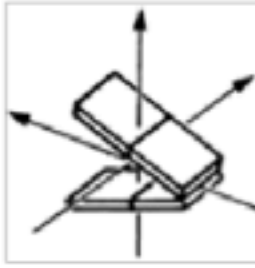

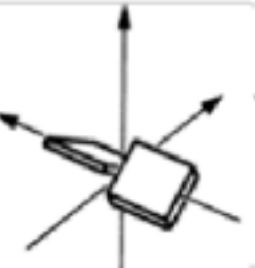
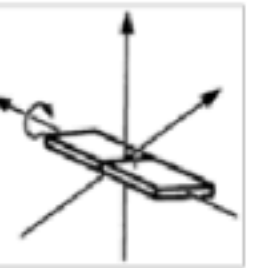
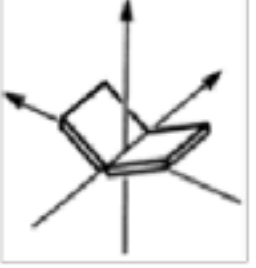
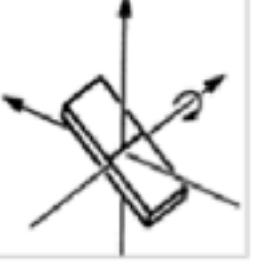
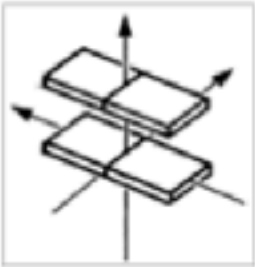
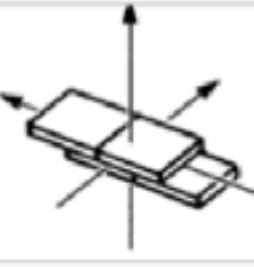
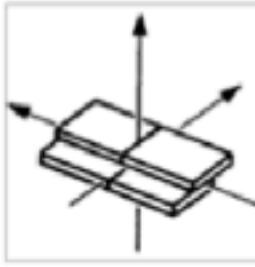
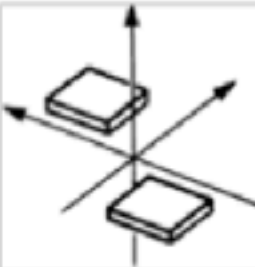
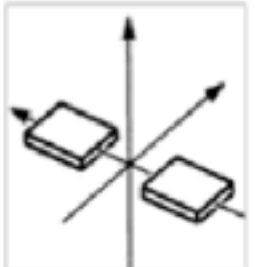
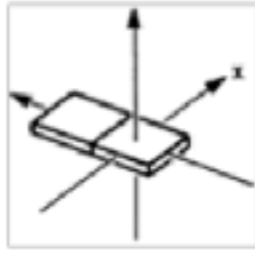
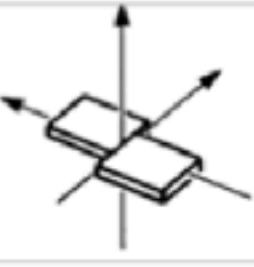
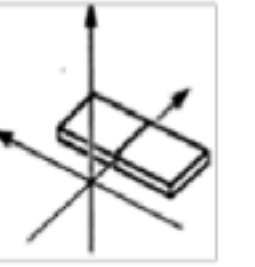
N. Seeman



DNA cube, Chen and Seeman. Nature 1991



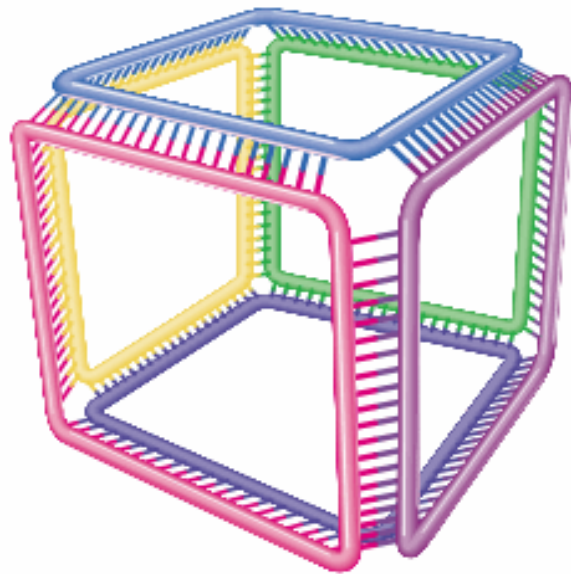
Geometrical parameters characterizing the structure of dsDNA

Base Step Parameters			Base Pair Parameters				
Rotational			Rotational				
							
Twist	Roll	Tilt	Opening	Propeller Twist	Tip	Buckle	Inclination
Translational			Translational				
							
Rise	Slide	Shift	Stagger	Stretch	Y displacement	Shear	X displacement
Z	Y	X	Z	Y	Y	X	X

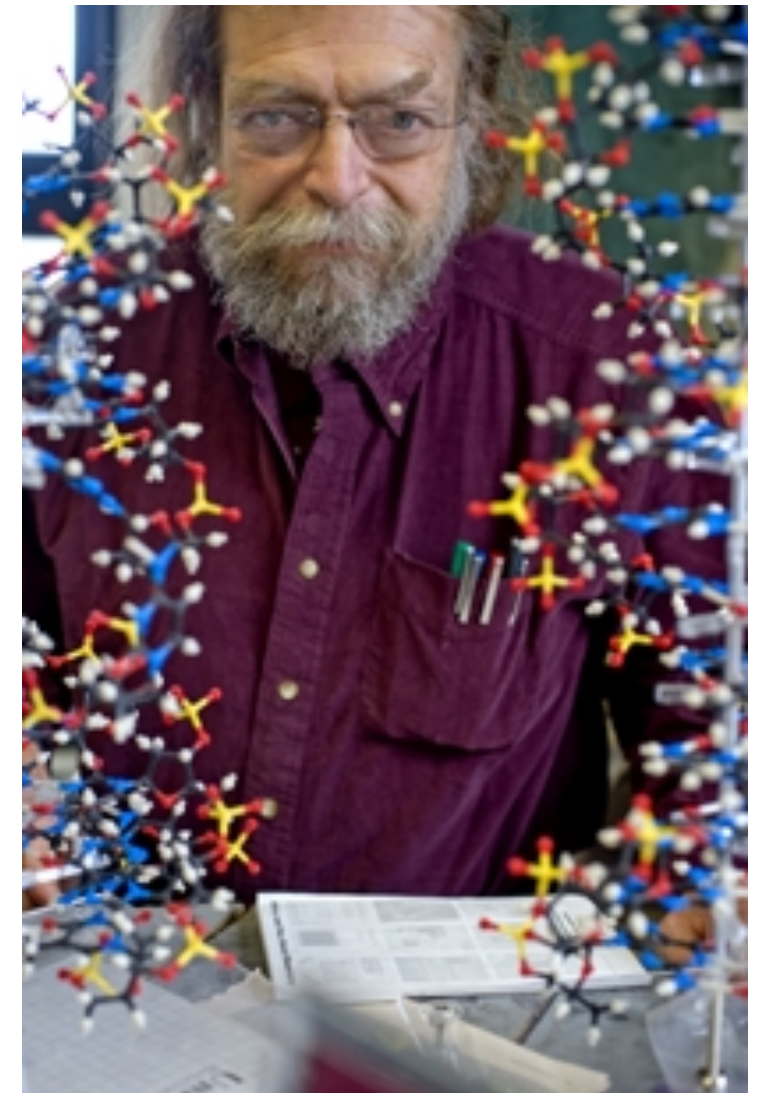
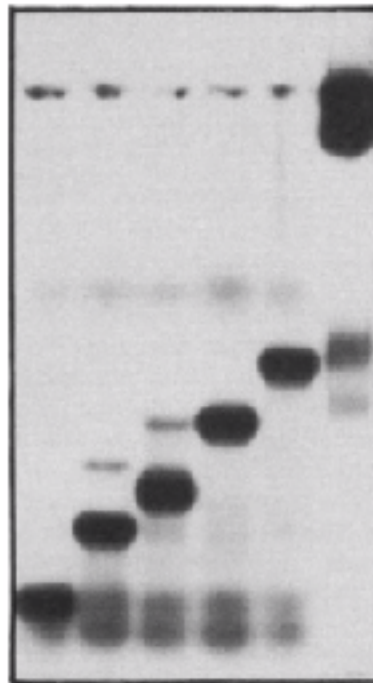
Geometrical parameters of various form of dsDNA

Structure	Helical Twist	Helical Rise	X-Displ	Inclin .	Roll	Propeller Twist	Diameter
A-DNA	33°	2.56 Å	-4.5 Å	21°	6°	-7.5°	23 Å
B-DNA	36°	3.38 Å	0.23 Å	-6°	0°	-4.4°	20 Å
Z-DNA	-30°	3.7Å	3.0 Å	-6.2°	0°	-4.4°	18 Å

DNA Nanotechnology



DNA cube,
Chen and Seeman. Nature 1991



DNA Nanotechnology

- A method to create nanoscale object using DNA/RNA as construction material
- Incepted by Nadrian C. Seeman, in 1982 to crystalize protein which is a very hard problem to solve.
- The field was inspired by using DNA nanostructure as a tool to in biophysics but now taken as a entire

Chemist saw it as a water-soluble polymer whose growth could be controlled and whose branching, or cross-linking, could be rationally manipulated.

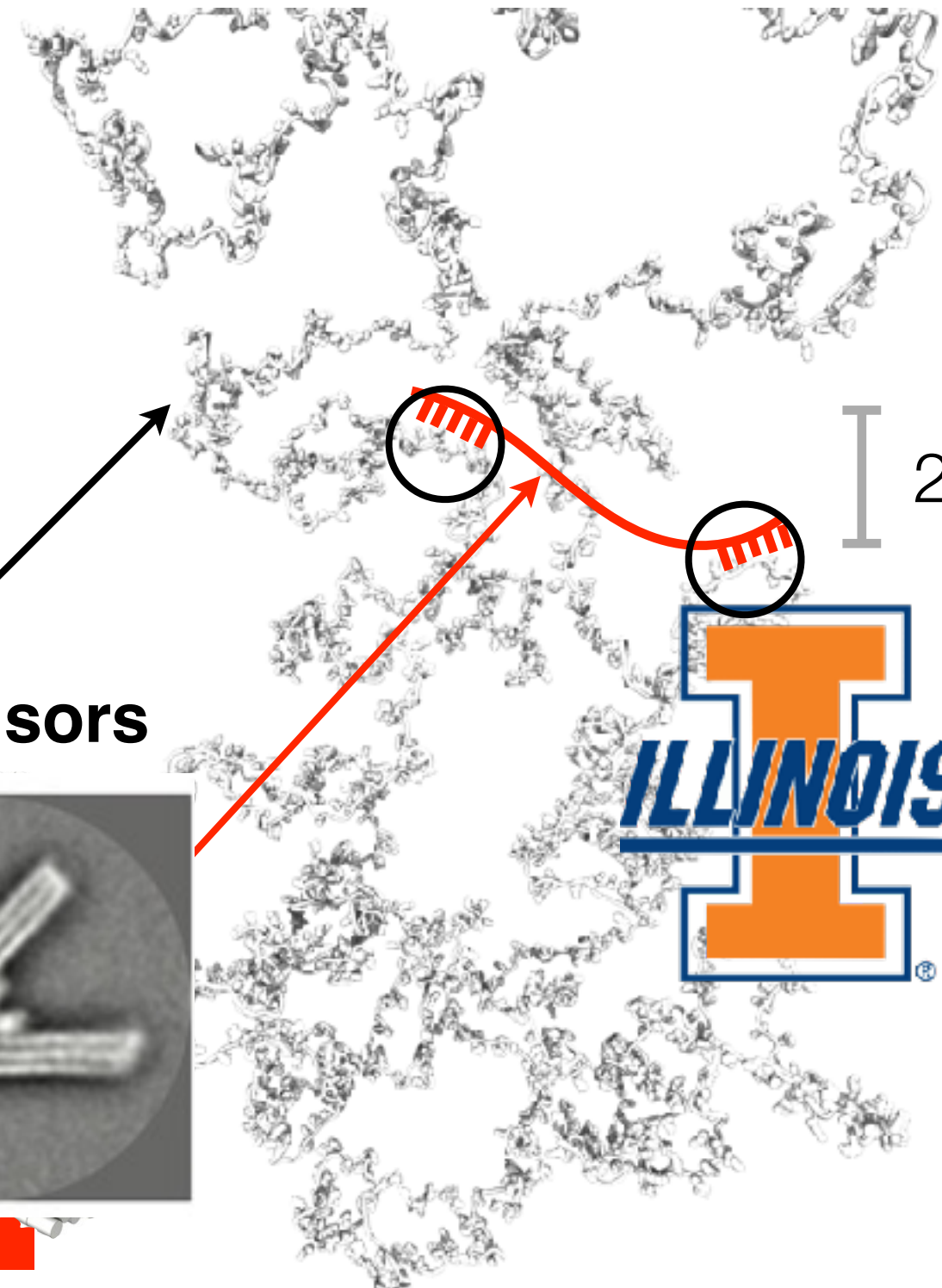
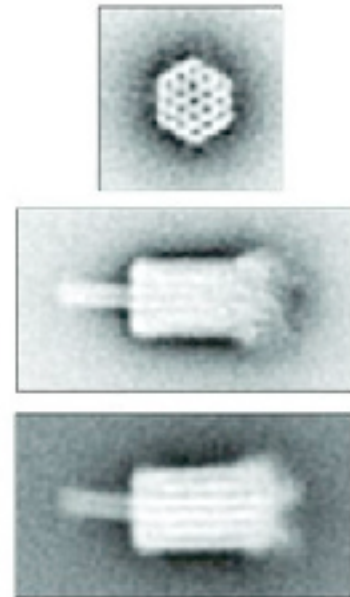
Yamuna and Ned, 2019



Ned Seeman
1945-2021

DNA Origami

Custom shapes, channels, and sensors



Nature 2005
Paul Rothemund


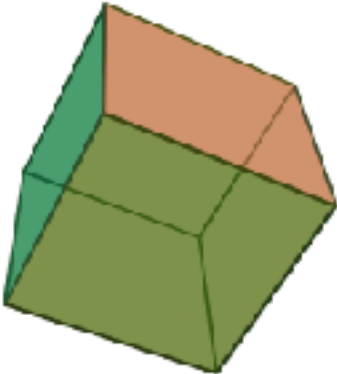
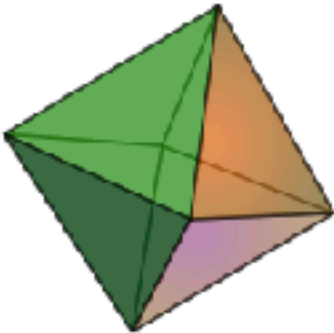
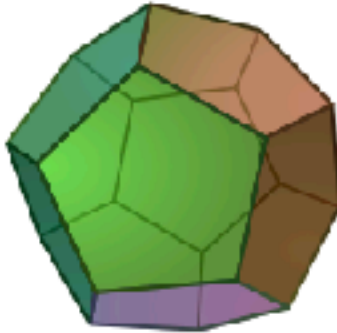
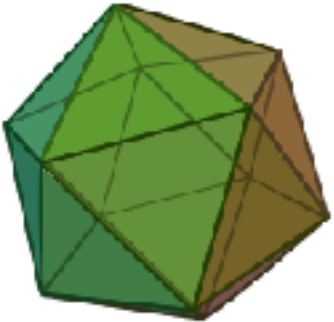
Science 2012
Simmel et al.

Science Adv. 2016
Dietz et al.

Source: J. Yoo, Aksimentiev lab

Platonic solids

Only five convex regular polyhedron exist in 3d euclidian space

Tetrahedron	Cube	Octahedron	Dodecahedron	Icosahedron
Four faces	Six faces	Eight faces	Twelve faces	Twenty faces
				

Parameters

Figure	Tetrahedron		Octahedron	Cube	Icosahedron		Dodecahedron	
Faces	4		8	6	20		12	
Vertices	4		6 (2 × 3)	8	12 (4 × 3)		20 (8 + 4 × 3)	
Position	1	2			1	2	1	2
Vertex coordinates	(1, 1, 1) (1, -1, -1) (-1, 1, -1) (-1, -1, 1)	(-1, -1, -1) (-1, 1, 1) (1, -1, 1) (1, 1, -1)	(±1, 0, 0) (0, ±1, 0) (0, 0, ±1)	(±1, ±1, ±1)	(0, ±1, ±φ) (±1, ±φ, 0) (±φ, 0, ±1)	(0, ±φ, ±1) (±φ, ±1, 0) (±1, 0, ±φ)	(± $\frac{1}{\phi}$, ±φ, 0) (±φ, 0, ± $\frac{1}{\phi}$)	(±φ, ± $\frac{1}{\phi}$, 0) (± $\frac{1}{\phi}$, 0, ±φ)

Virus assembly

Using VMD

Download VMD

<https://www.ks.uiuc.edu/Training/Tutorials/vmd/tutorial-html/node1.html>

Visualizing a cube in VMD

Tcl programming

Usage: Visualization, Building models, Analysis of simulation Trajectories

Homework

Laws of thermodynamics

DNA translational and rotational parameters

Reading material