ME 411 / ME 511

## Biological Frameworks for Engineers





## Class Organization

• HW1 assigned. Due Wed Oct 3.



ME 411 / ME 511

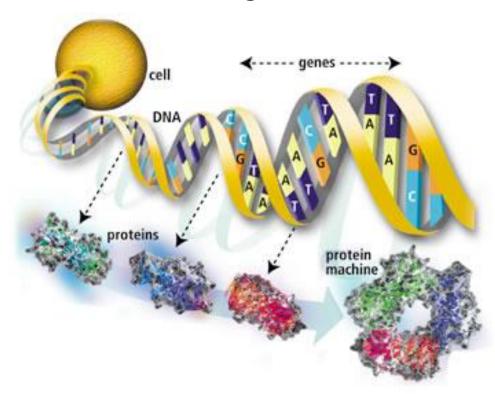
## Information Handling





## Deoxyribonucleic Acid

DNA is a polymer of nucleotides that encodes the sequence of amino acid in proteins through a template called messenger RNA

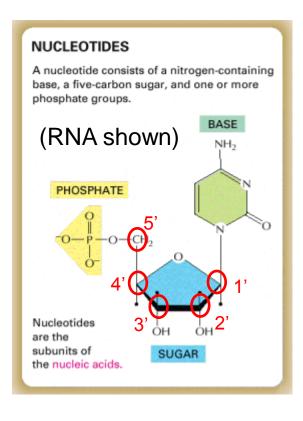






## Nucleotides

Consists of a nitrogen base, 5-carbon sugar, and a phosphate from the 5' end of the sugar.



### <u>Sugar</u>

Ribonucleic acid (RNA)
Deoxyribonucleic acid (DNA)

## Nitrogen Bases

G: Guanine (Purine)
A: Adenine (Purine)
T: Thymine (Pyrimidine)
C: Cytosine (Pyrimidine)
U: Uracil (Pyrimidine)

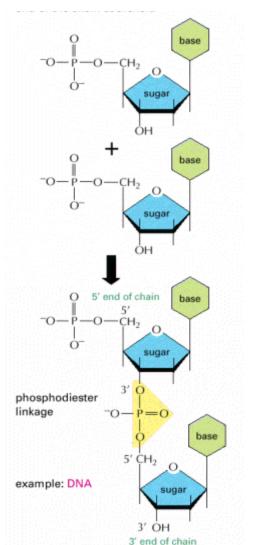




## DNA Polymerization

A nucleic acid strand has a pentosephosphate backbone.

A phosphodiester bond connects the 3' end of one nucleic acid with the 5' end of the next mer.



(DNA shown)

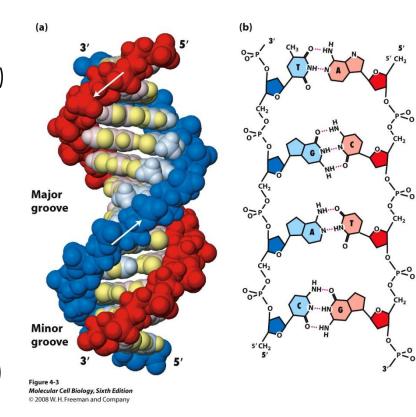


## cal Frameworks for **Engineers**

## DNA Structure

1953: Watson and Crick proposed the double-helix structure of DNA\*

- Right-handed, Anti-parallel
  - 5' / 3'  $\rightarrow$  3' / 5' (Strands run opposite)
- Base pairs on the inside
  - A T has two H-bonds
  - G C has three H-bonds
- Dimensions
  - 2.0 nm wide
  - 3.4 nm length per turn
  - 10 base pairs per turn
- Grooves
  - Major: 2.2 nm wide (protein docking)
  - Minor: 1.2 nm wide

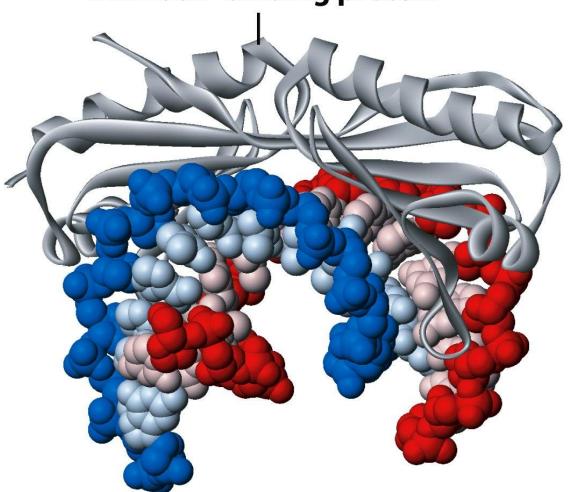




<sup>\*</sup> using Maurice Wilkins and Rosalind Franklin's X-ray diffraction data.

## It's Bend-y!

## **TATA box-binding protein**



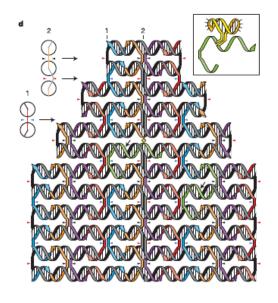
Flexibility arises because H-bonds run perpendicular and not parallel to long axis





# gical Frameworks for Engineers

## It's Constructable!



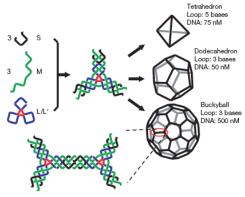
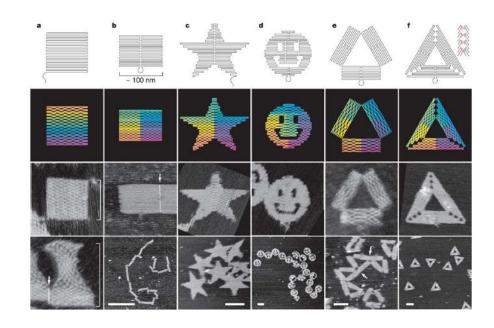
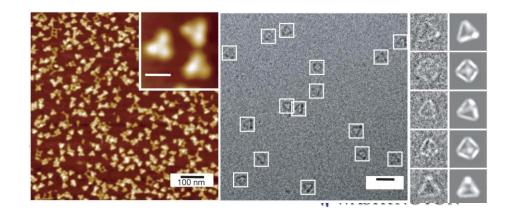


Figure 1 | Self-assembly of DNA polyhedra. Three different types of DNA single strands stepwise assemble into symmetric three-point-star motifs (tiles) and then into polyhedra in a one-pot process. There are three single-stranded loops (coloured red) in the centre of the complex. The final structures (polyhedra) are determined by the loop length (3 or 5 bases long) and the DNA concentration.





## Hw 1...

Explore information handling in Replication (DNA → DNA)

Transcription (DNA → RNA) & Translation (RNA → Proteins)

