

Nucleic Acids & Replication

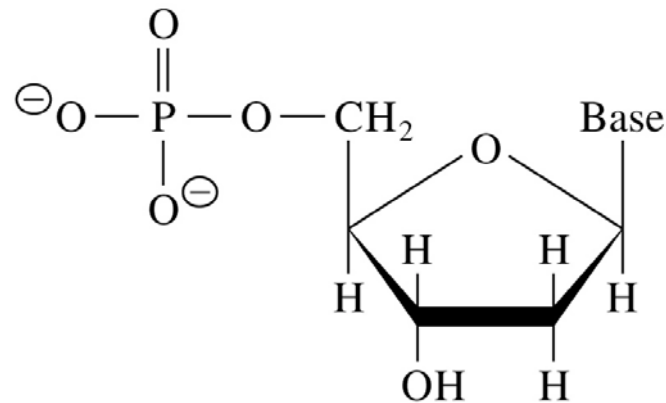
- **Nucleic acids** represent the fourth major class of biomolecules (other major classes of biomolecules are proteins, carbohydrates, fats)
- **Genome** - the genetic information of an organism

Information flow:

DNA → **RNA** → **PROTEIN**

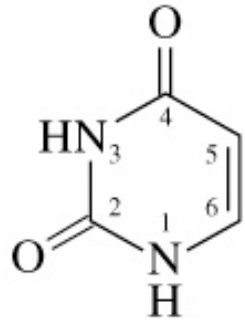
Nucleotides Are the Building Blocks of Nucleic Acids

- Nucleic acids are polynucleotides
- Nucleotides are phosphate esters of **nucleosides**



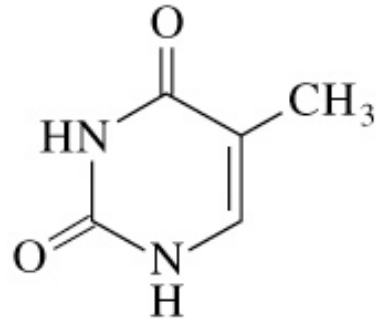
Major pyrimidines and purines

PYRIMIDINES



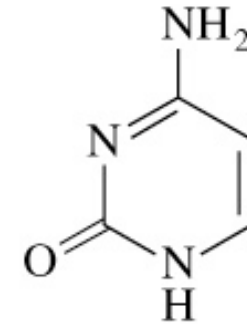
Uracil

(2,4-Dioxypyrimidine)



Thymine

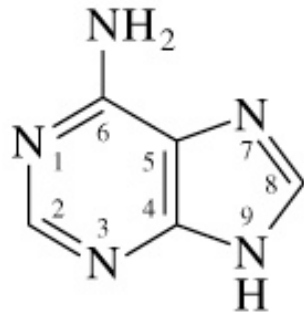
(2,4-Dioxo-5-methylpyrimidine)



Cytosine

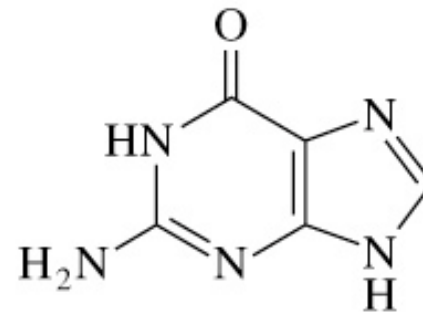
(2-Oxo-4-aminopyrimidine)

PURINES



Adenine

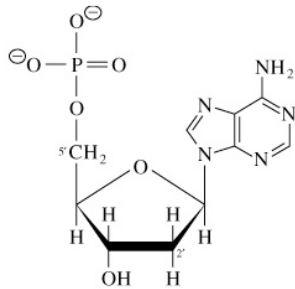
(6-Aminopurine)



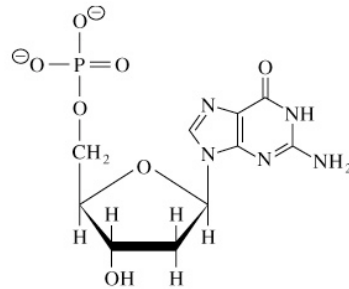
Guanine

(2-Amino-6-oxopurine)

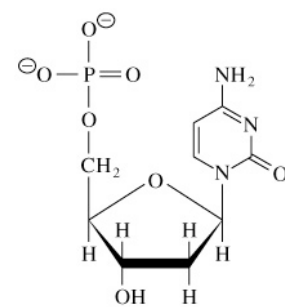
Structures of the deoxyribonucleoside-5'-monophosphates



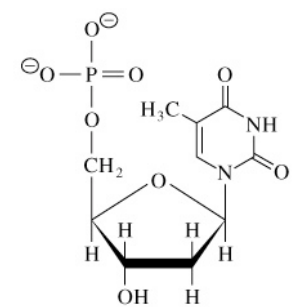
2'-Deoxyadenosine 5'-monophosphate
(Deoxyadenylate, dAMP)



2'-Deoxyguanosine 5'-monophosphate
(Deoxyguanylate, dGMP)

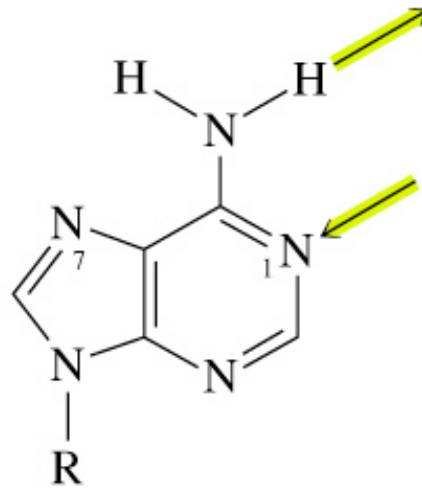


2'-Deoxycytidine 5'-monophosphate
(Deoxycytidylate, dCMP)

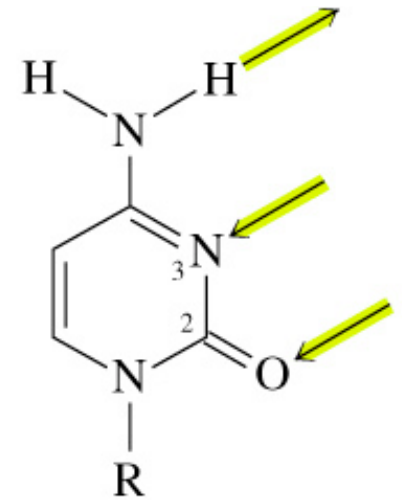


2'-Deoxythymidine 5'-monophosphate
(Thymidylate, dTMP)

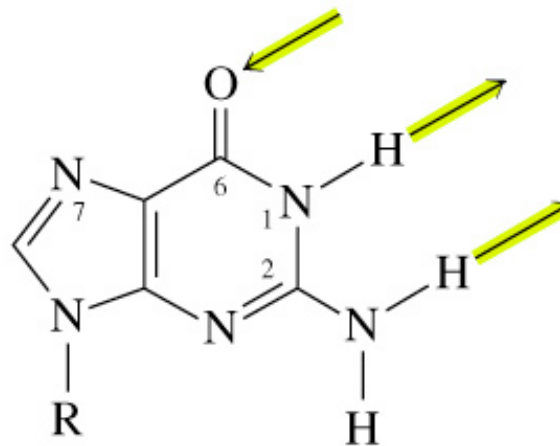
- Hydrogen bond sites in nucleic acids



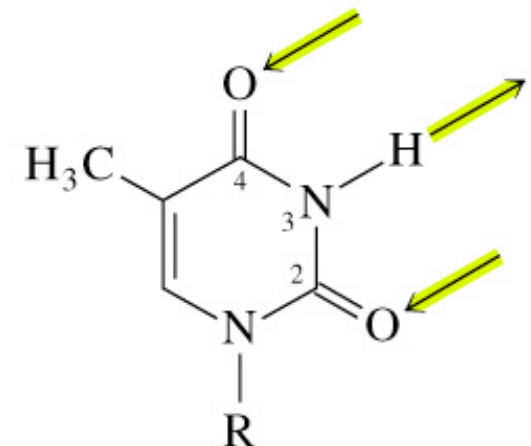
(Deoxy)Adenosine



(Deoxy)Cytidine



(Deoxy)Guanosine

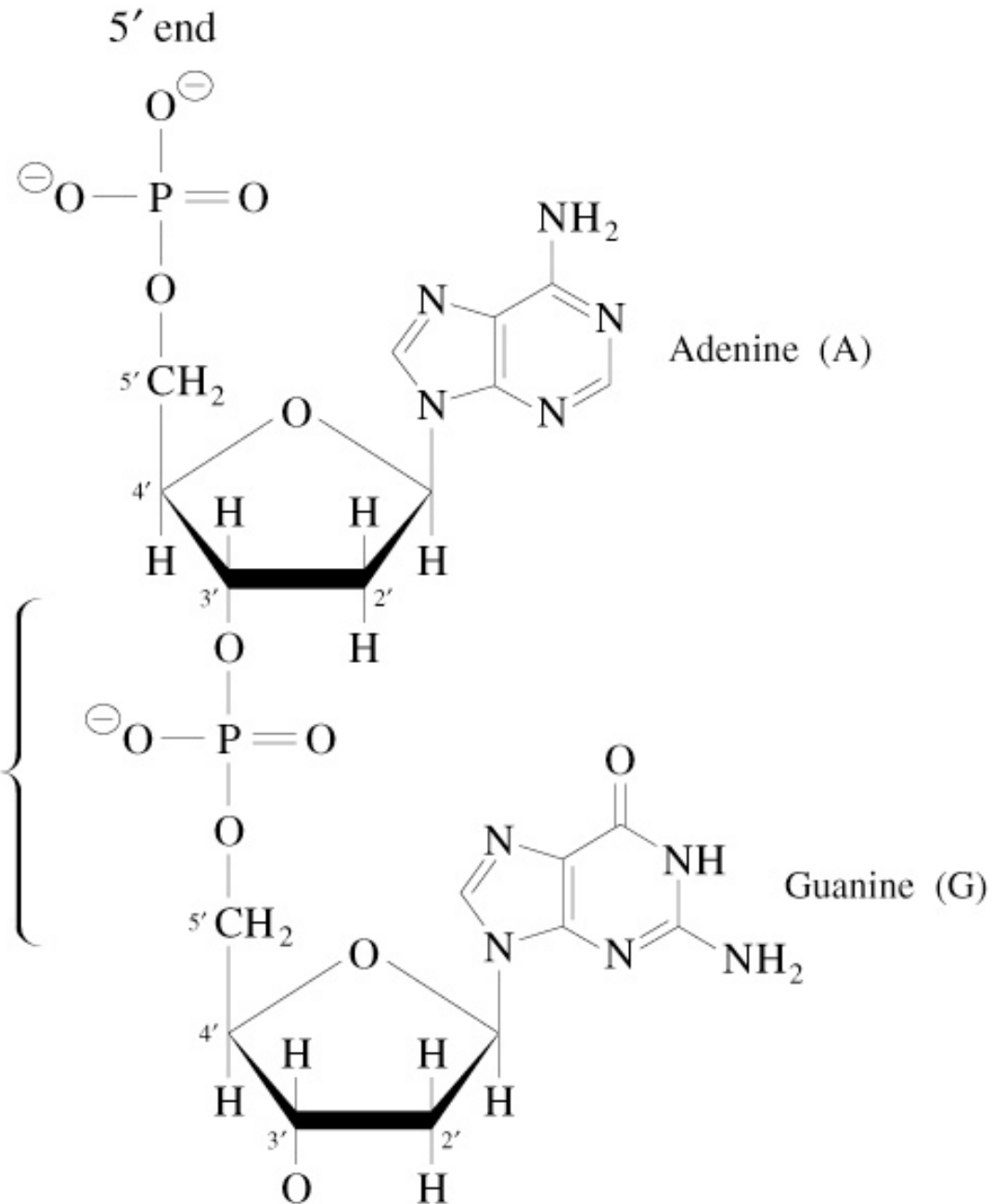


(Deoxy)Thymidine

Nucleotides joined by 3'-5' phosphodiester linkages

Structure of the
tetranucleotide
pdApdGpdTpdC

3'-5'
phosphodiester
linkage



19.2 DNA Is Double-Stranded

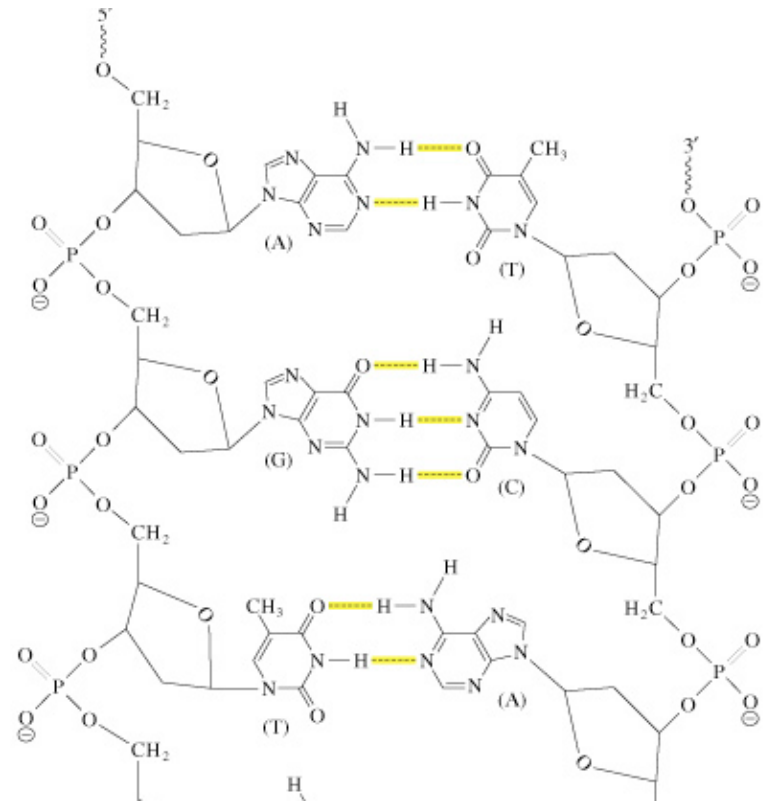
Table 19.2

| TABLE 19.2 Base composition of DNA(mole%)and ratios of bases | | | | | | | | |
|--|------|------|------|------|------------------|------------------|---------|------------------------------------|
| Source | A | G | C | T | A/T ^a | G/C ^a | (G + C) | Purine/ pyrimidine ^a |
| <i>Escherichia coli</i> | 26.0 | 24.9 | 25.2 | 23.9 | 1.09 | 0.99 | 50.1 | 1.04 |
| <i>Mycobacterium tuberculosis</i> | 15.1 | 34.9 | 35.4 | 14.6 | 1.03 | 0.99 | 70.3 | 1.00 |
| Yeast | 31.7 | 18.3 | 17.4 | 32.6 | 0.97 | 1.05 | 35.7 | 1.00 |
| Cow | 29.0 | 21.2 | 21.2 | 28.7 | 1.01 | 1.00 | 42.4 | 1.01 |
| Pig | 29.8 | 20.7 | 20.7 | 29.1 | 1.02 | 1.00 | 41.4 | 1.01 |
| Human | 30.4 | 19.9 | 19.9 | 30.1 | 1.01 | 1.00 | 39.8 | 1.01 |

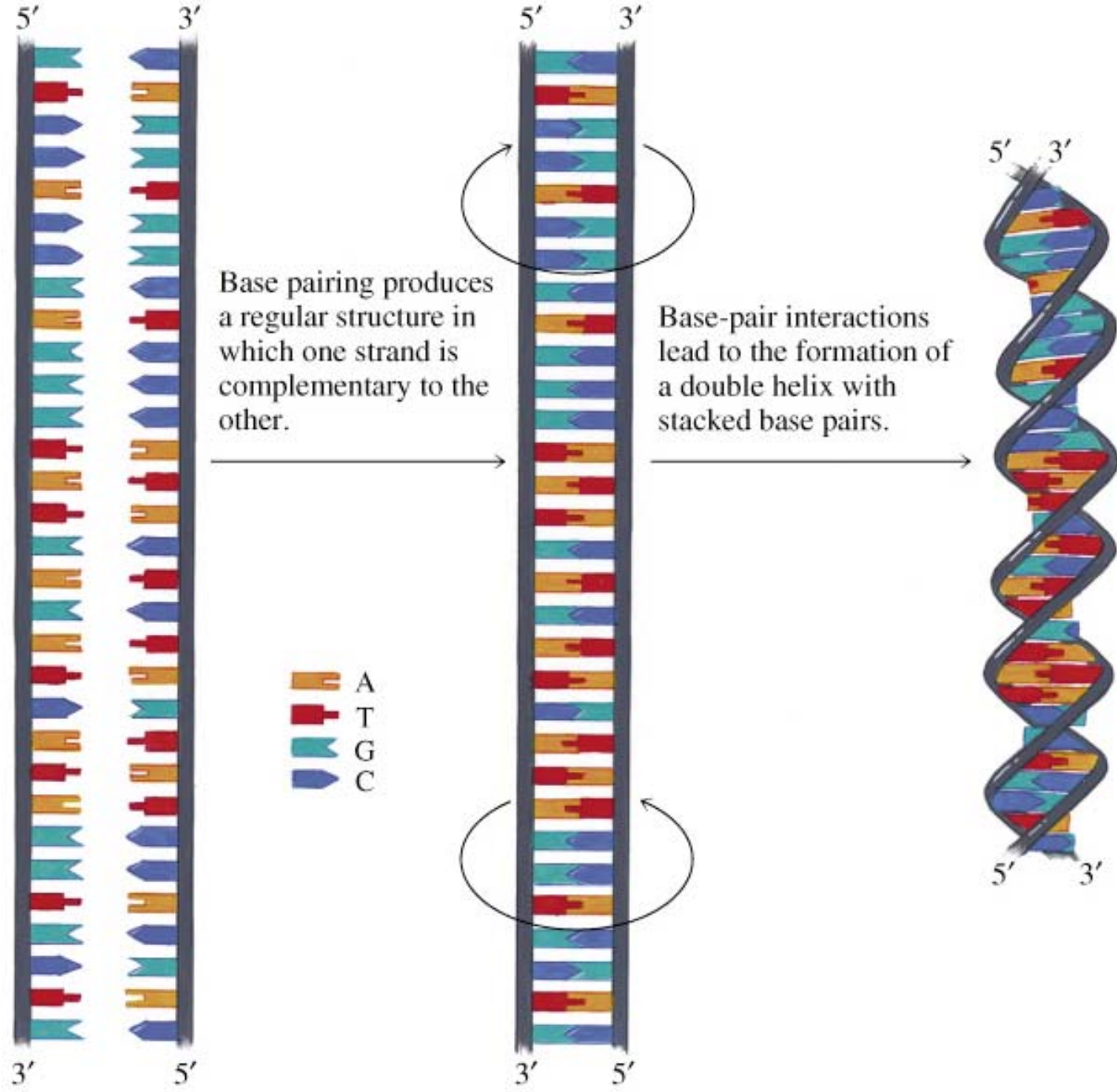
^aDeviations from a 1:1 ratio are due to experimental variations.

Two Antiparallel Strands Form a Double Helix

- Two strands run in opposite directions
- Bases in opposite strands pair by complementary hydrogen bonding
- Adenine (A) - Thymine (T)
- Guanine (G) - Cytosine (C)



- Complementary base pairing and stacking in DNA

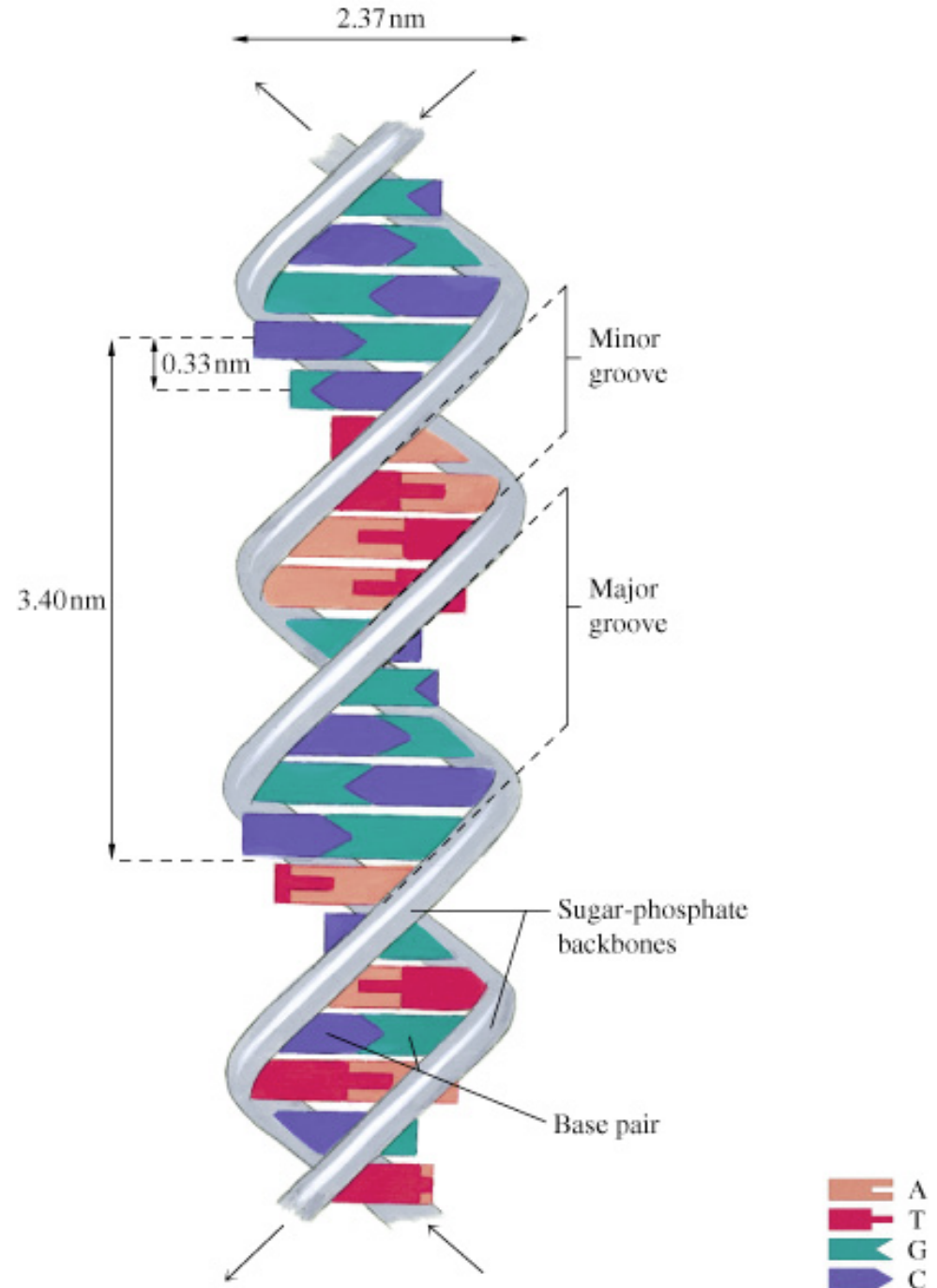


Three dimensional structure of DNA

- A double helix has two grooves of unequal width: **major groove** and **minor groove**
- Within each groove base pairs are exposed and are accessible to interactions with other molecules
- DNA-binding proteins can use these interactions to “read” a specific sequence

Fig 19.14

- Structure of B-DNA
- Sugar phosphate backbone outside
- Stacking creates two unequal grooves (major and minor)



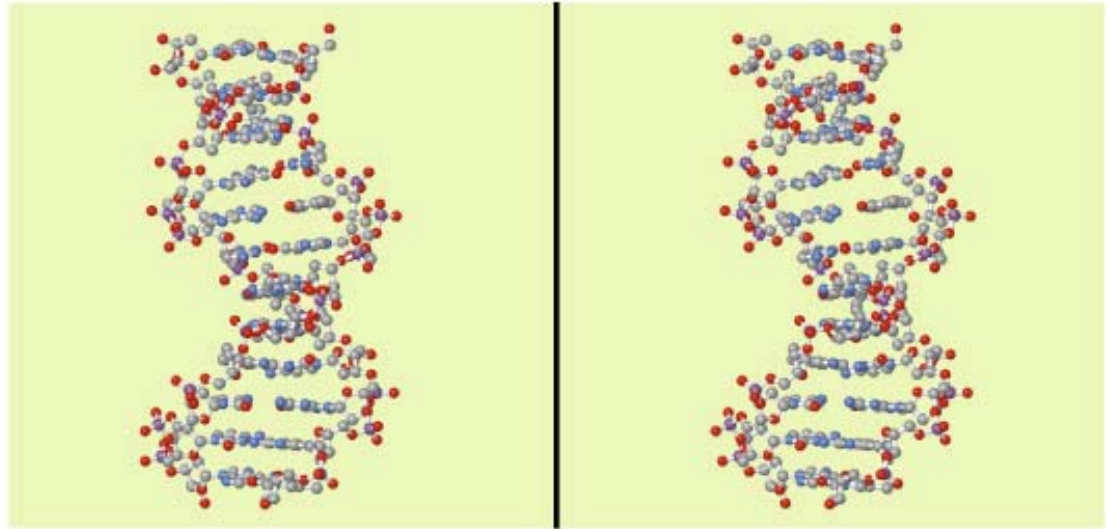
Stereo view of B-DNA

B-DNA is a right-handed helix, diam. = 2.37nm

- **Rise** (distance between stacked bases) = 0.33nm
- **Pitch** (distance to complete one turn) = 3.40 nm
- Base pairs nearly perpendicular to sugar-phosphate backbones
- **Figure 19.15** Stereo views of B-DNA (next slide)

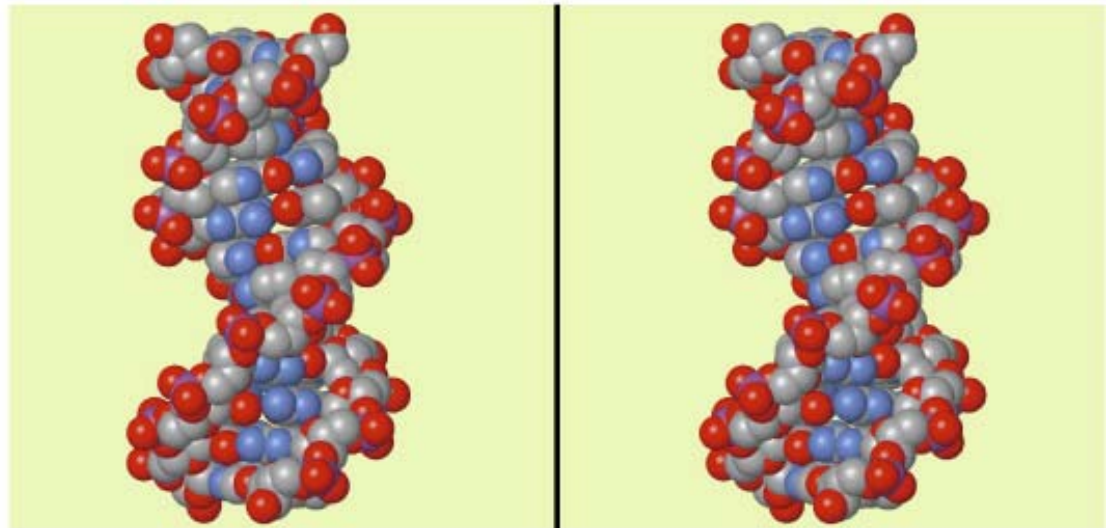
(a) Ball-and-stick model

(a)



(b) Space-filling model

(b)



Weak Forces Stabilize the Double Helix

- (1) *Hydrophobic effects.* Burying purine and pyrimidine rings in the double helix interior
- (2) *Stacking interactions.* Stacked base pairs form van der Waals contacts

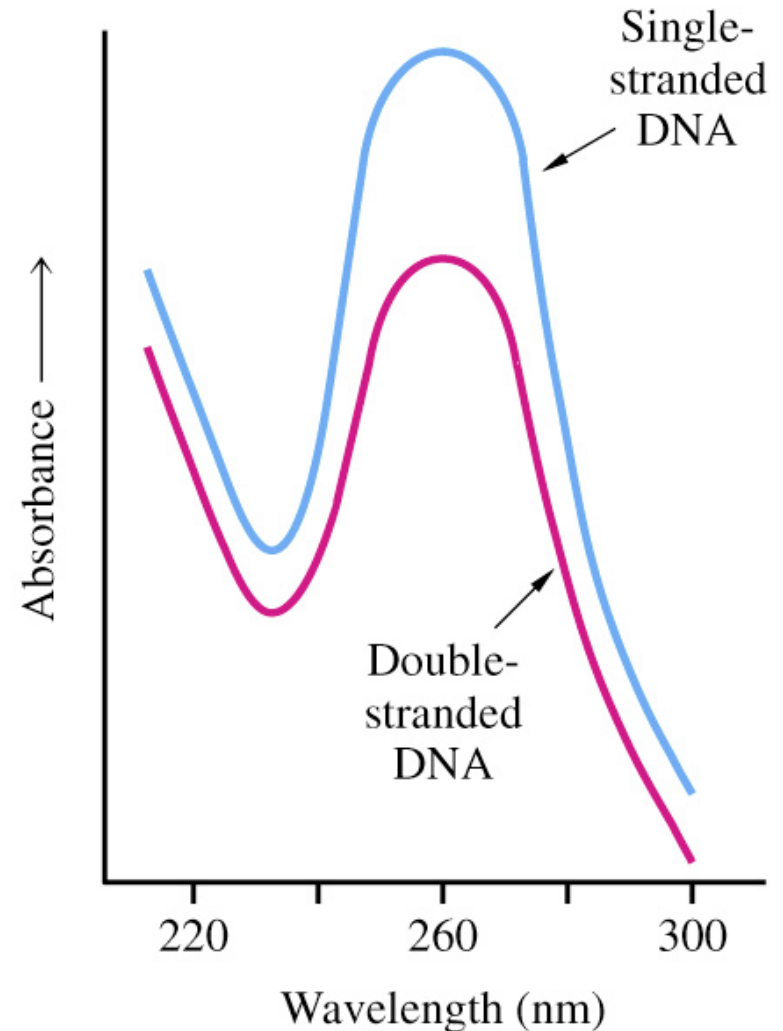
(continued next slide)

Weak forces (continued)

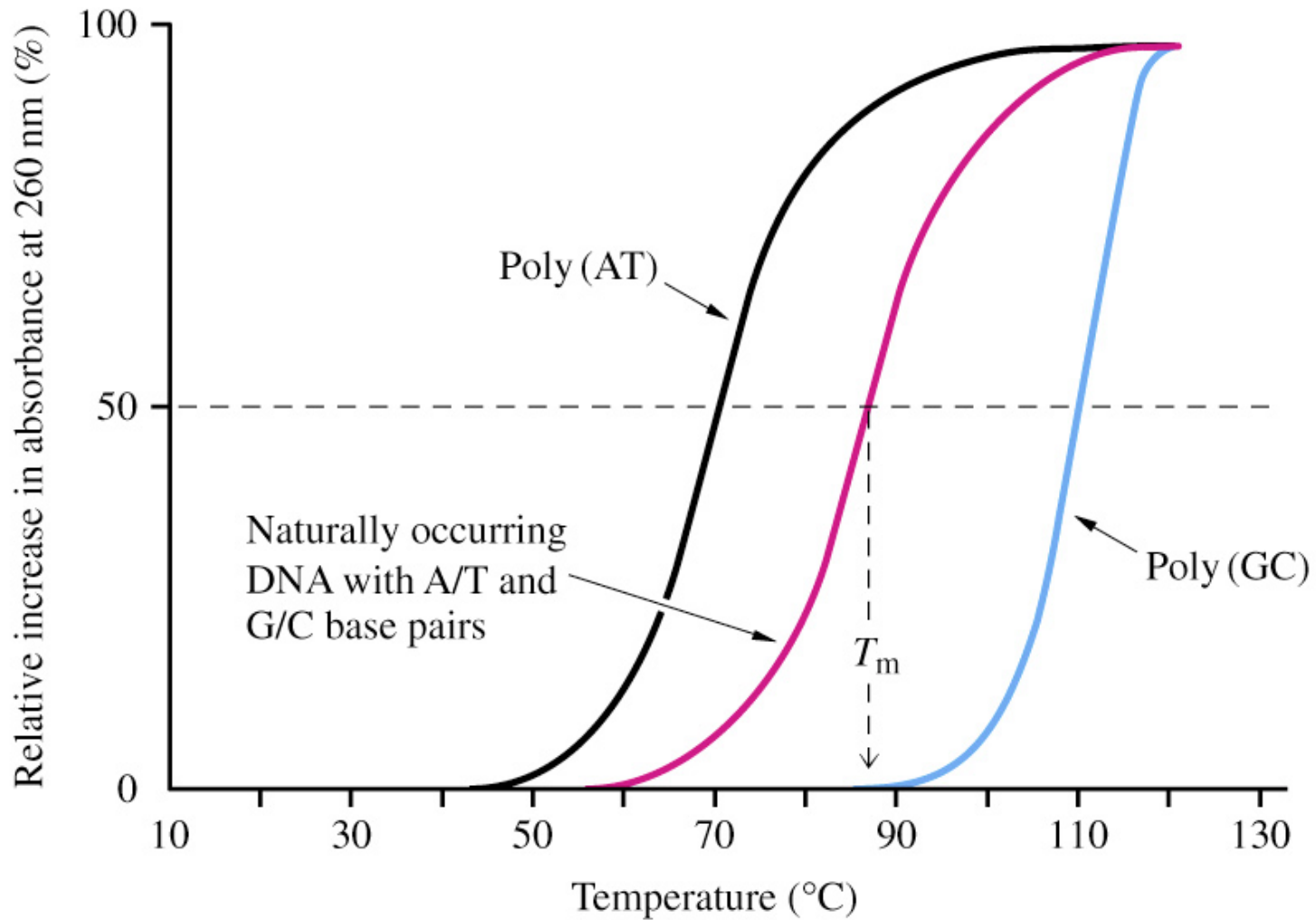
- (3) *Hydrogen bonds.* Hydrogen bonding between base pairs.
- (4) *Charge-charge interactions.* Electrostatic repulsion of negatively charged phosphate groups is decreased by cations (e.g. Mg^{2+}) and cationic proteins

Absorption spectra of double-stranded and single-stranded DNA

- Double-stranded (DS)DNA (pH 7.0), absorbance max 260nm
- Denatured DNA absorbs 12% -40% more than DS DNA



Melting curve for DNA

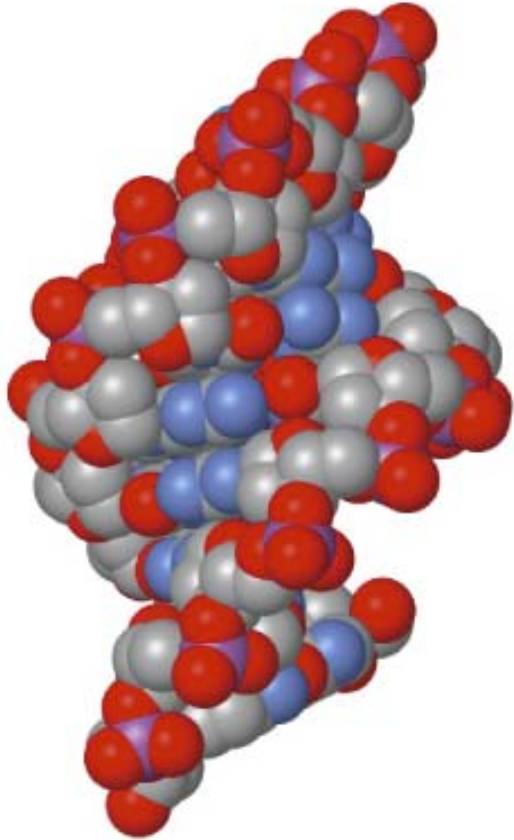


Conformations of Double-Stranded DNA

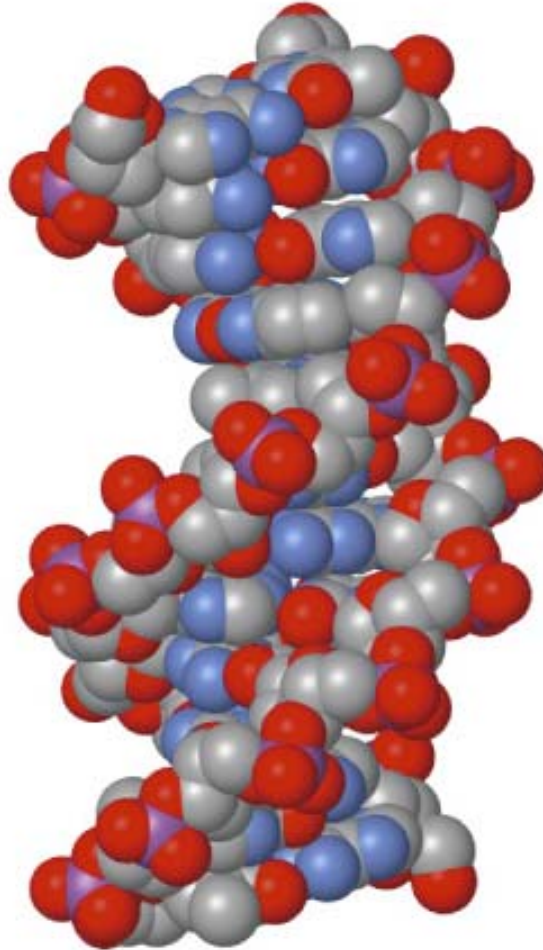
- Two alternative structures to B-DNA:
 - A-DNA** (forms when DNA is dehydrated)
 - Z-DNA** (when certain sequences are present)
- A-DNA is more tightly wound than B-DNA, and has minor grooves of similar width
- Z-DNA has no grooves and a left-handed helix
- Both A-DNA and Z-DNA exist in vivo in short regions of DNA

Forms of DNA

A



B



Z

