



Molecular Biology (1)

DNA structure and basic applications

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Second semester, 2018-2019

Resources

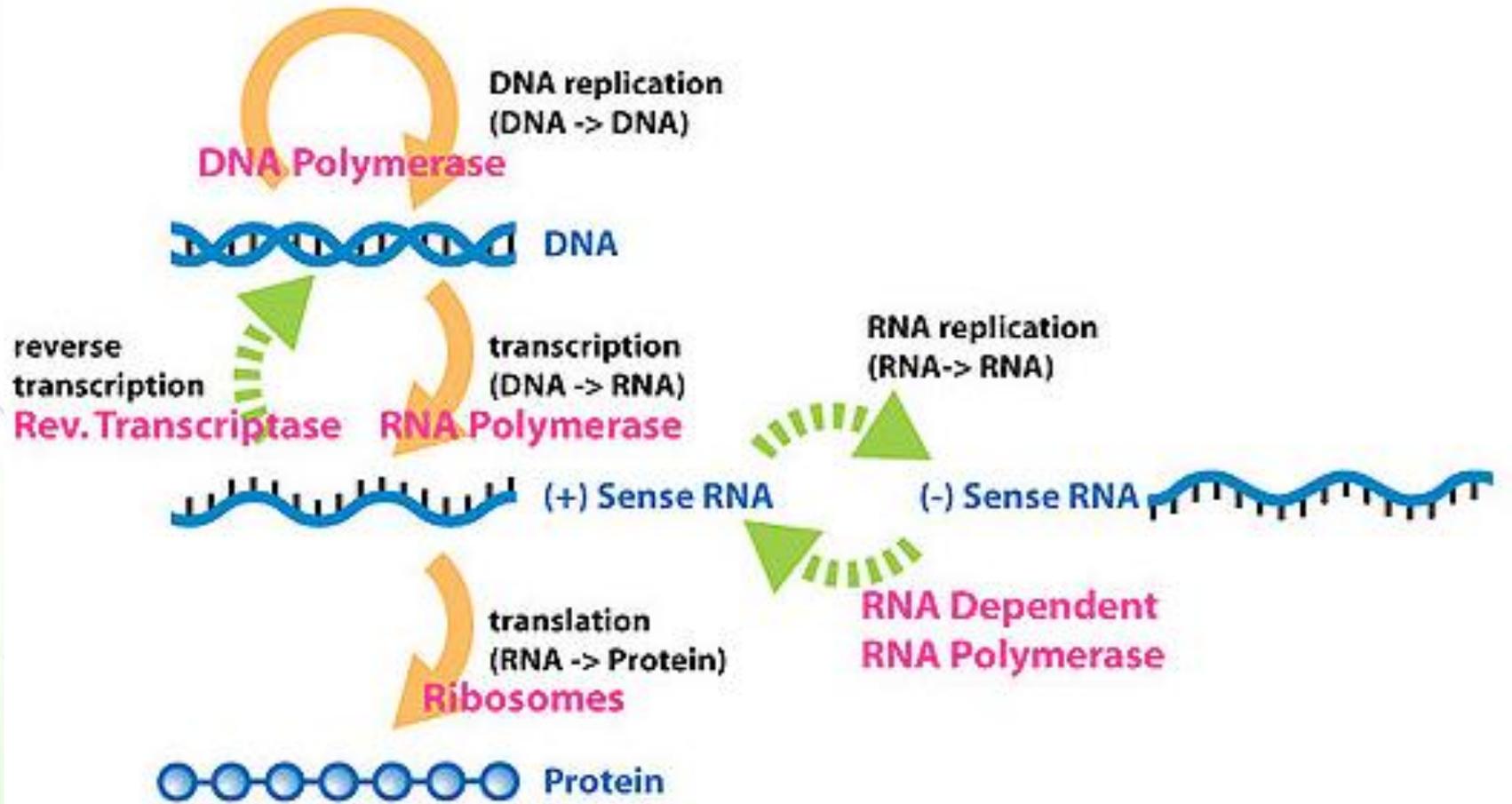


- This lecture
- Cooper, pp. 49-52, 118-119, 130

What is molecular biology?



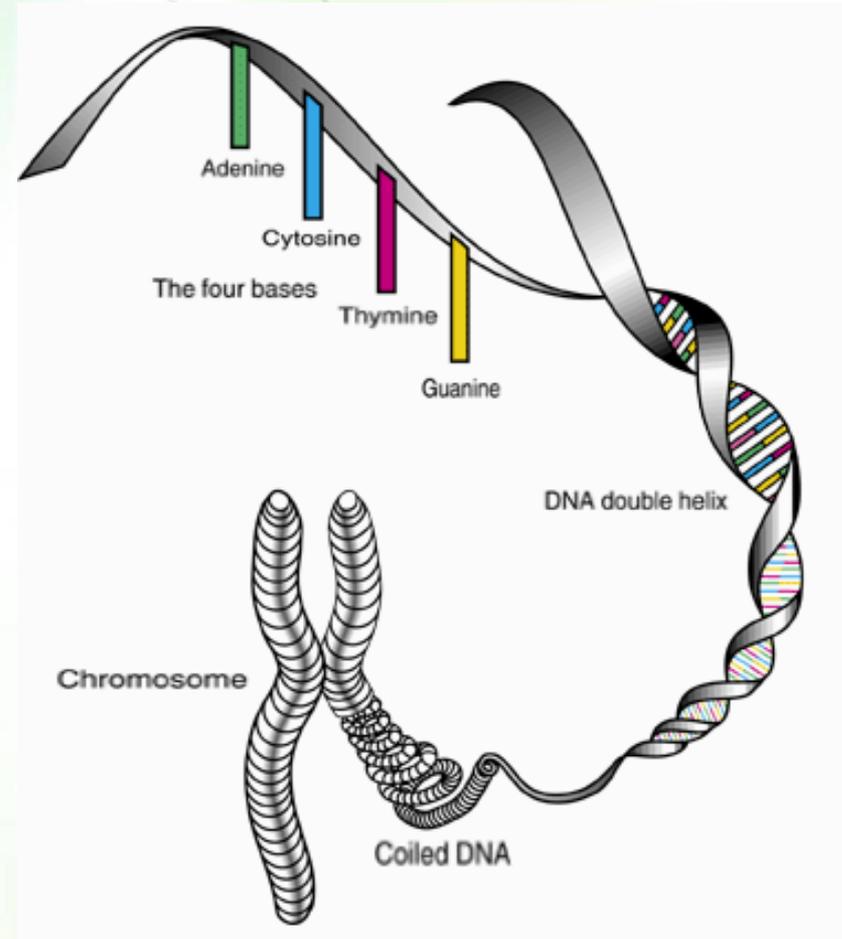
Central dogma of molecular biology



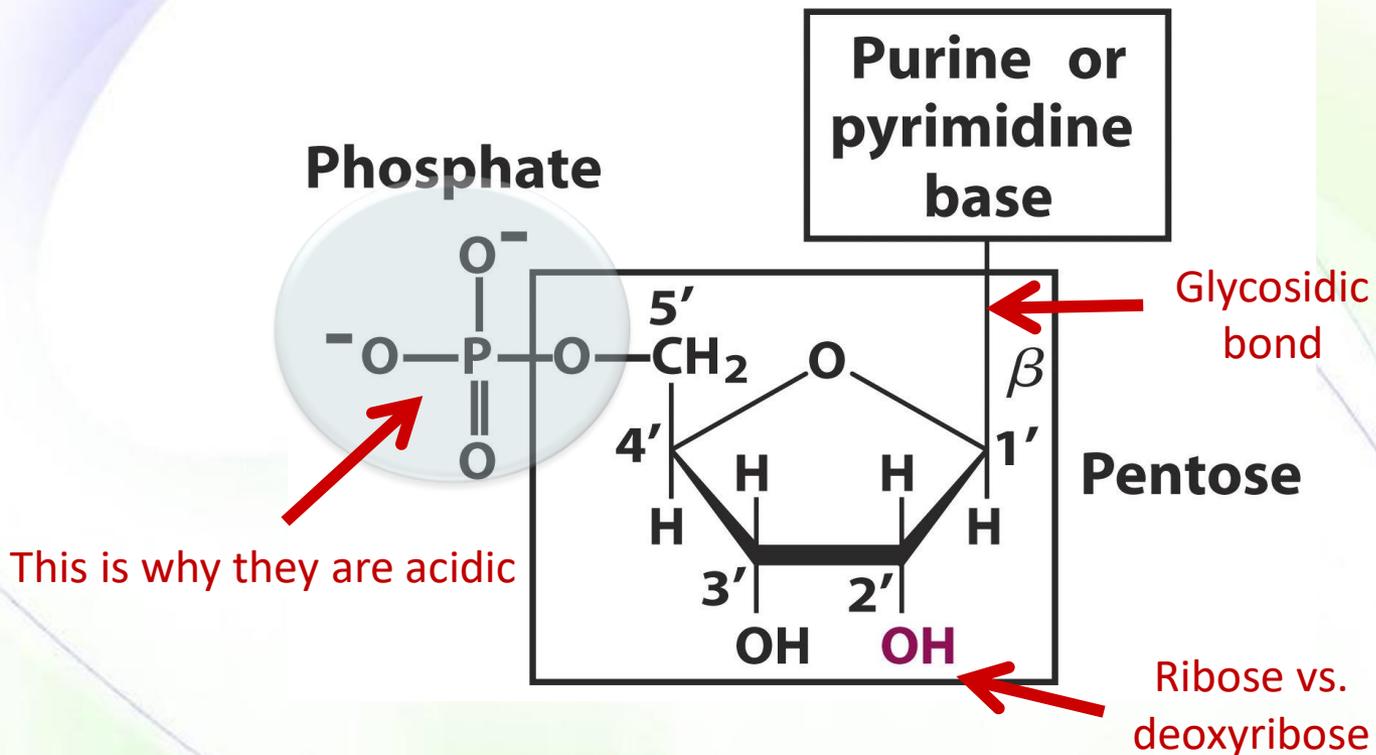
Nucleic acids



- The primary structure of nucleic acids is linear polymers of nucleotides (monomers) bound to each other via phosphodiester bonds.
- DNA is coiled and can be associated with proteins forming chromosomes.

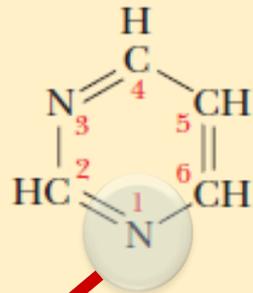


Chemical composition and bonds

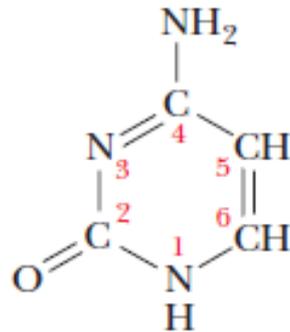


- Positively charged ions (Na^+ or Mg^{2+}) and peptides with positively charged side chains can associate with DNA
- Eukaryotic DNA, for example, is complexed with histones, which are positively charged proteins, in the cell nucleus.

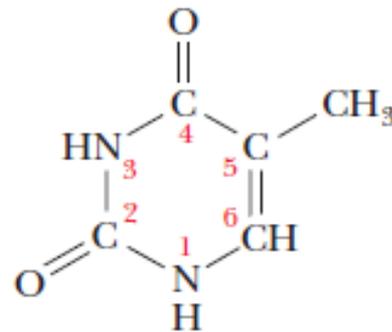
Nitrogenous bases



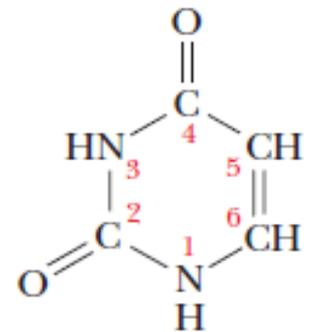
Pyrimidine



Cytosine
(in DNA & RNA)

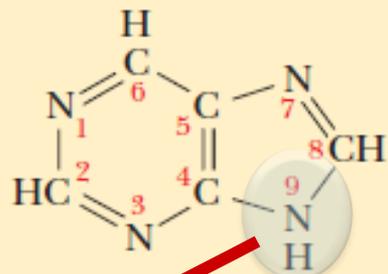


Thymine
(in DNA & some RNA)

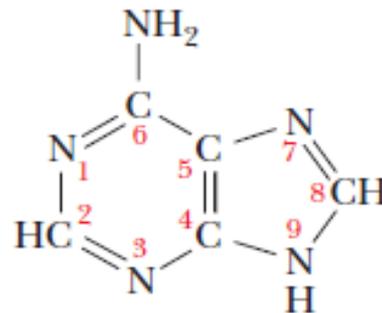


Uracil
(in RNA)

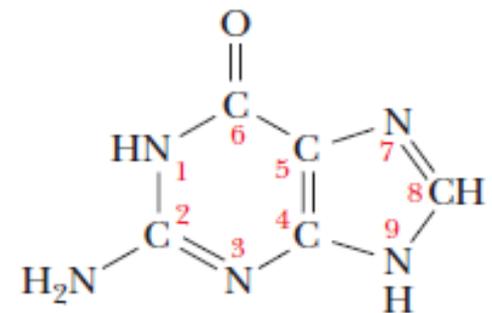
Glycosidic bond



Purine

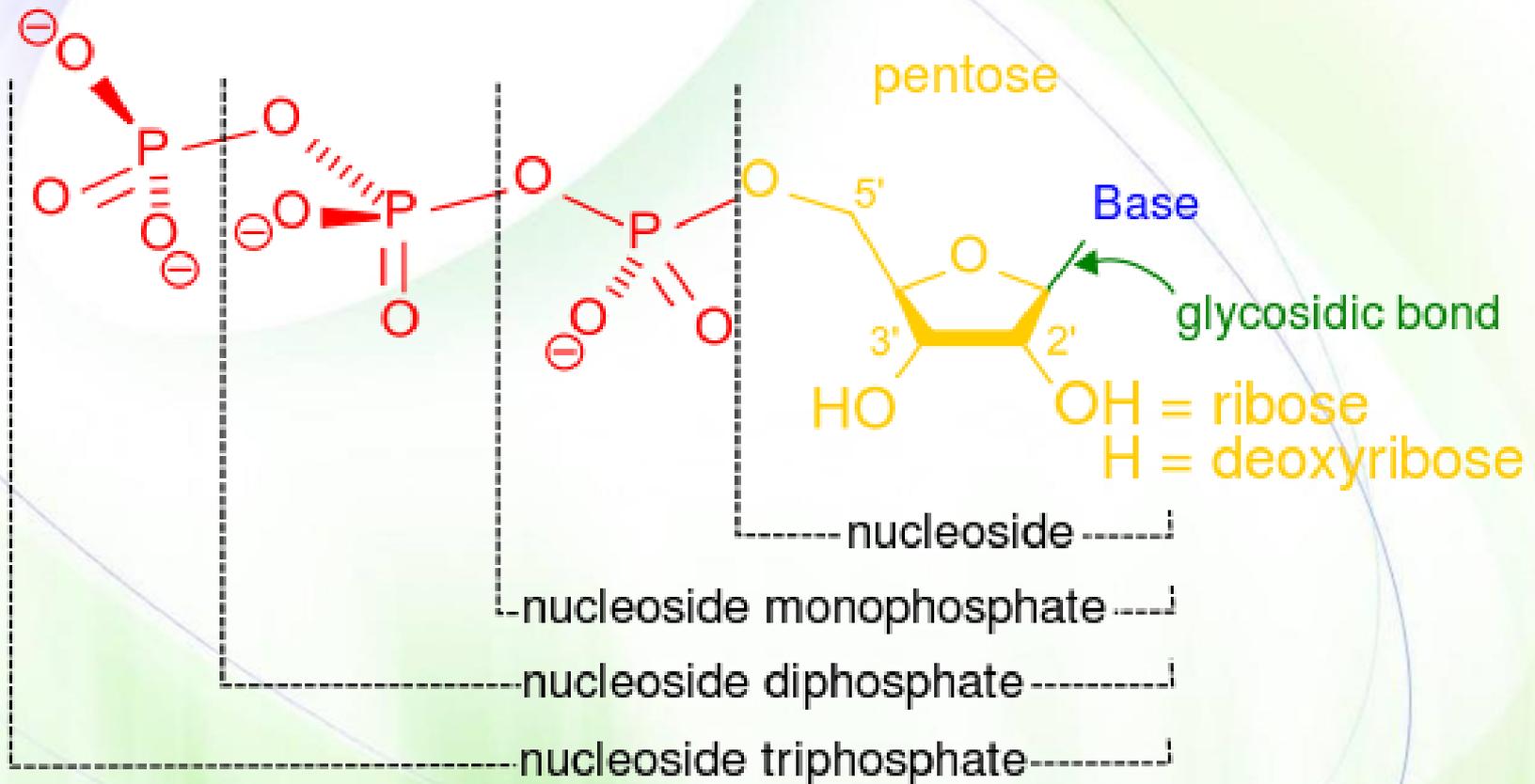


Adenine
(in DNA & RNA)

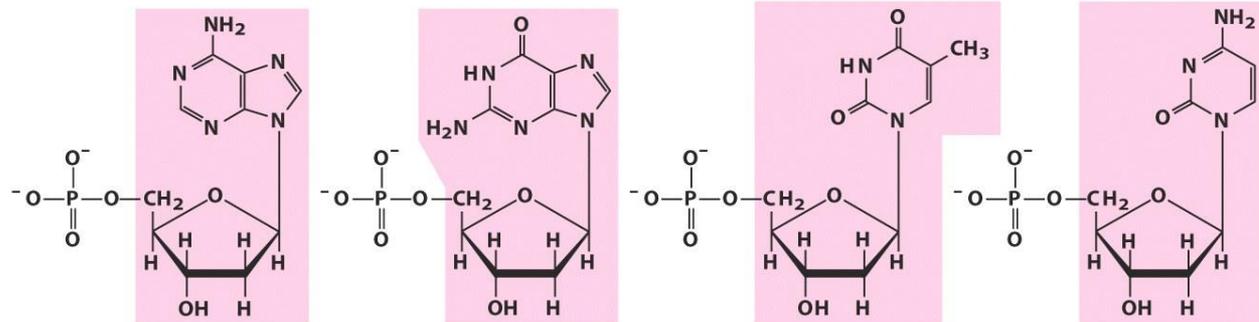


Guanine
(in DNA & RNA)

Nucleotides vs. Nucleosides



Nucleotides vs. Nucleosides



Nucleotide: Deoxyadenylate
(deoxyadenosine
5'-monophosphate)

Symbols: A, dA, dAMP

Nucleoside: Deoxyadenosine

Nucleotide: Deoxyguanylate
(deoxyguanosine
5'-monophosphate)

Symbols: G, dG, dGMP

Nucleoside: Deoxyguanosine

Nucleotide: Deoxythymidylate
(deoxythymidine
5'-monophosphate)

Symbols: T, dT, dTMP

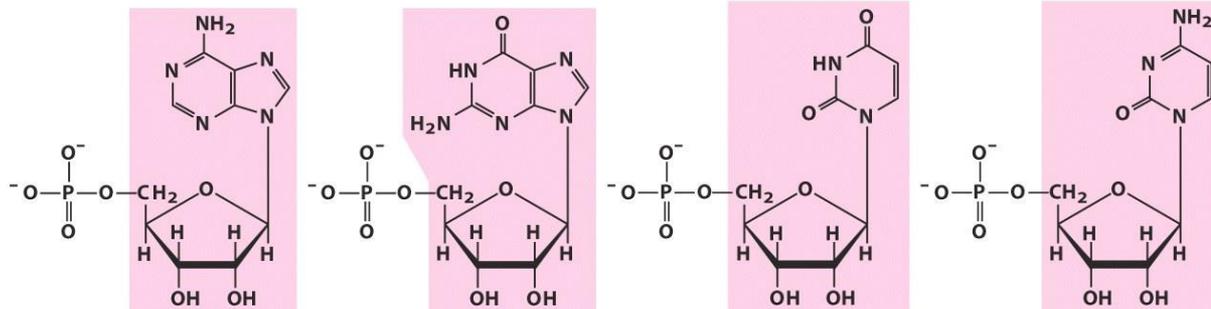
Nucleoside: Deoxythymidine

Nucleotide: Deoxycytidylate
(deoxycytidine
5'-monophosphate)

Symbols: C, dC, dCMP

Nucleoside: Deoxycytidine

(a) Deoxyribonucleotides



Nucleotide: Adenylate (adenosine
5'-monophosphate)

Symbols: A, AMP

Nucleoside: Adenosine

Nucleotide: Guanylate (guanosine
5'-monophosphate)

Symbols: G, GMP

Nucleoside: Guanosine

Nucleotide: Uridylate (uridine
5'-monophosphate)

Symbols: U, UMP

Nucleoside: Uridine

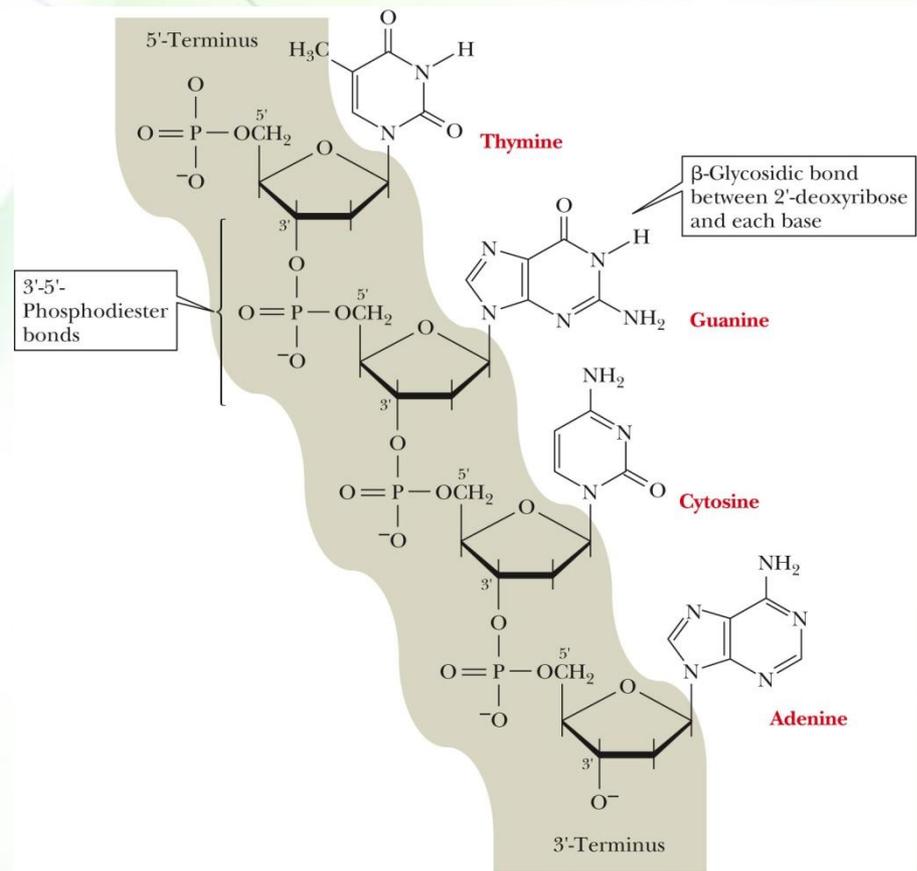
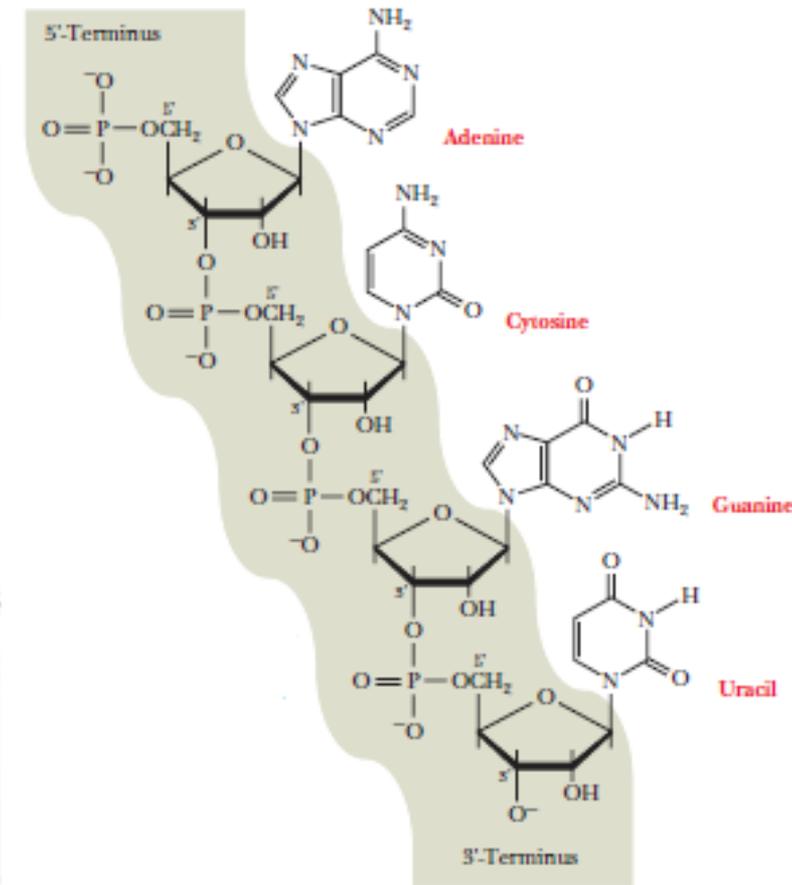
Nucleotide: Cytidylate (cytidine
5'-monophosphate)

Symbols: C, CMP

Nucleoside: Cytidine

(b) Ribonucleotides

Nucleic acid polymers

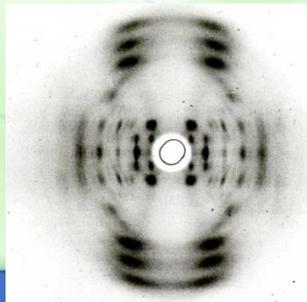
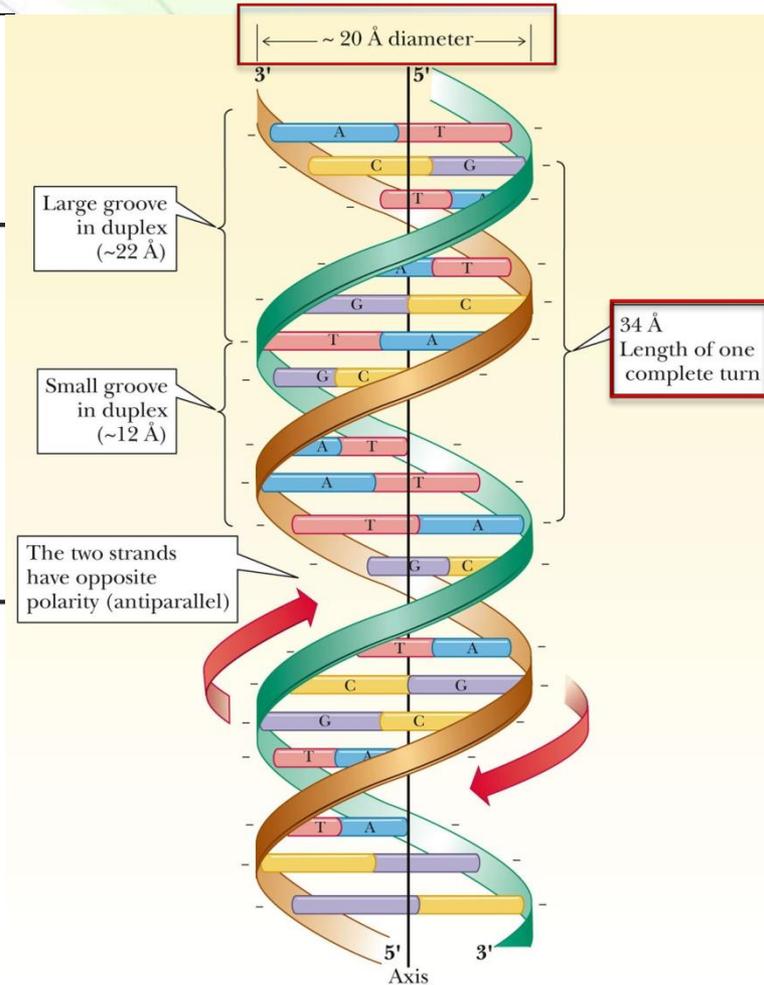
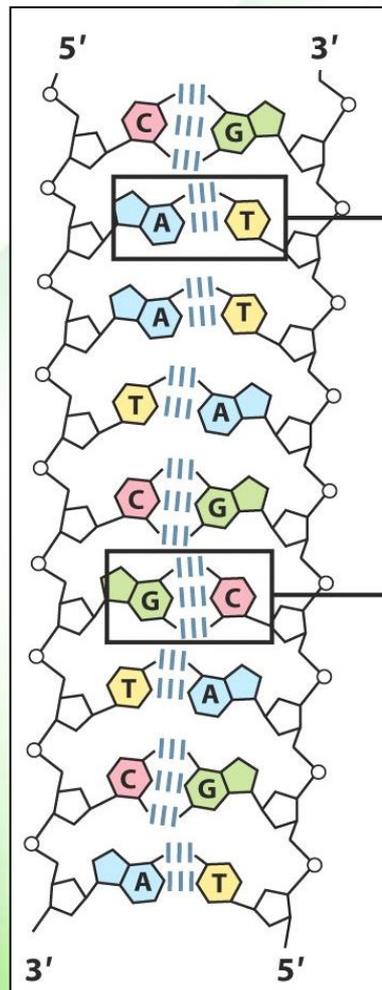


- A letter d can be added to indicate a deoxyribonucleotide residue.
- for example, dG is substituted for G.
- The deoxy analogue of a ribooligonucleotide would be d(GACAT).

DNA structure



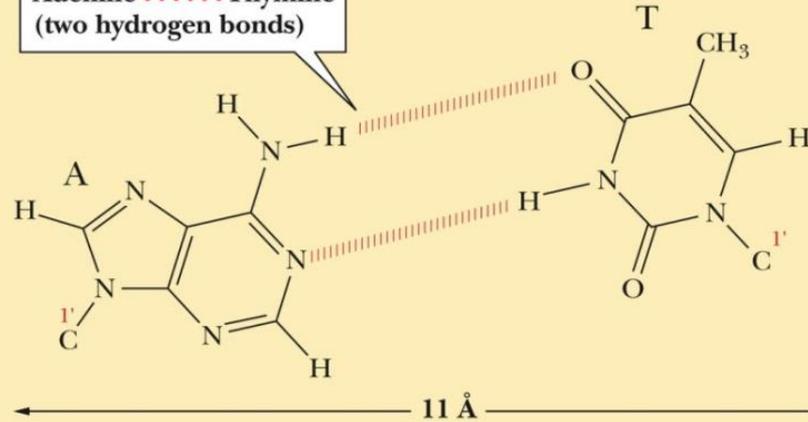
- Specific base-pairing
 - A = T; G = C; Pur = pyr
- Complementary
- A double helix
- Backbone vs. side chains
- Antiparallel
- Stable
- Flexible
- Groovings
- Stability vs. flexibility



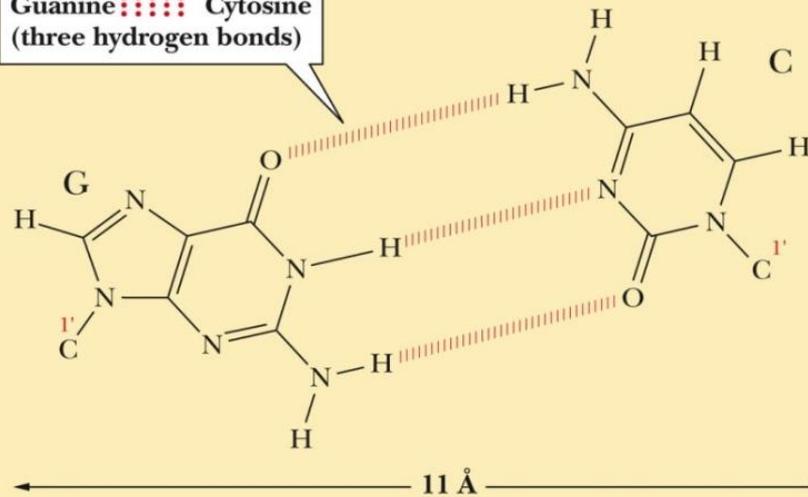
Base pairing



Adenine :::::Thymine
(two hydrogen bonds)



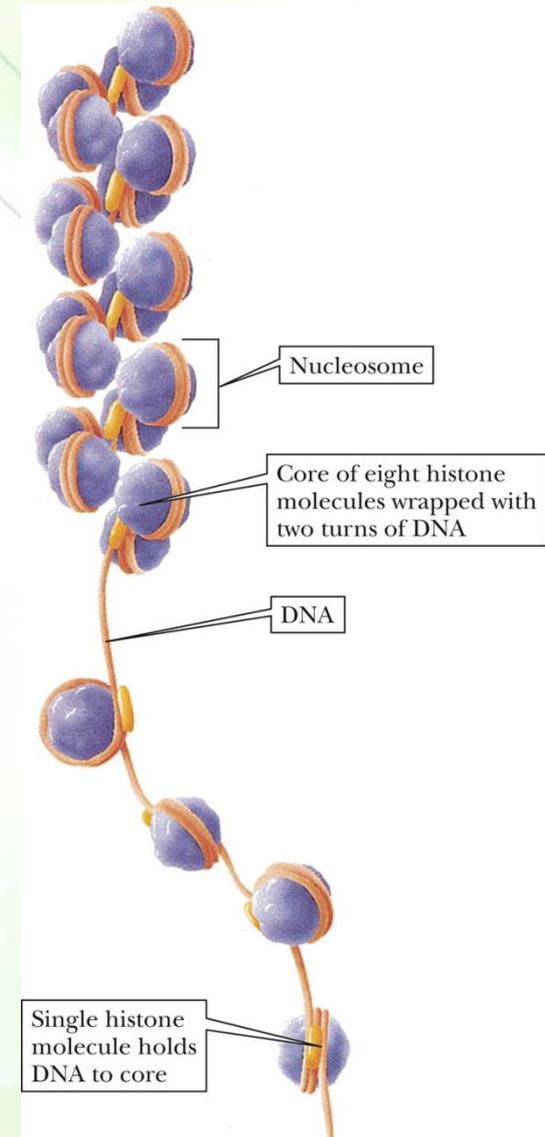
Guanine ::::: Cytosine
(three hydrogen bonds)



In eukaryotes...

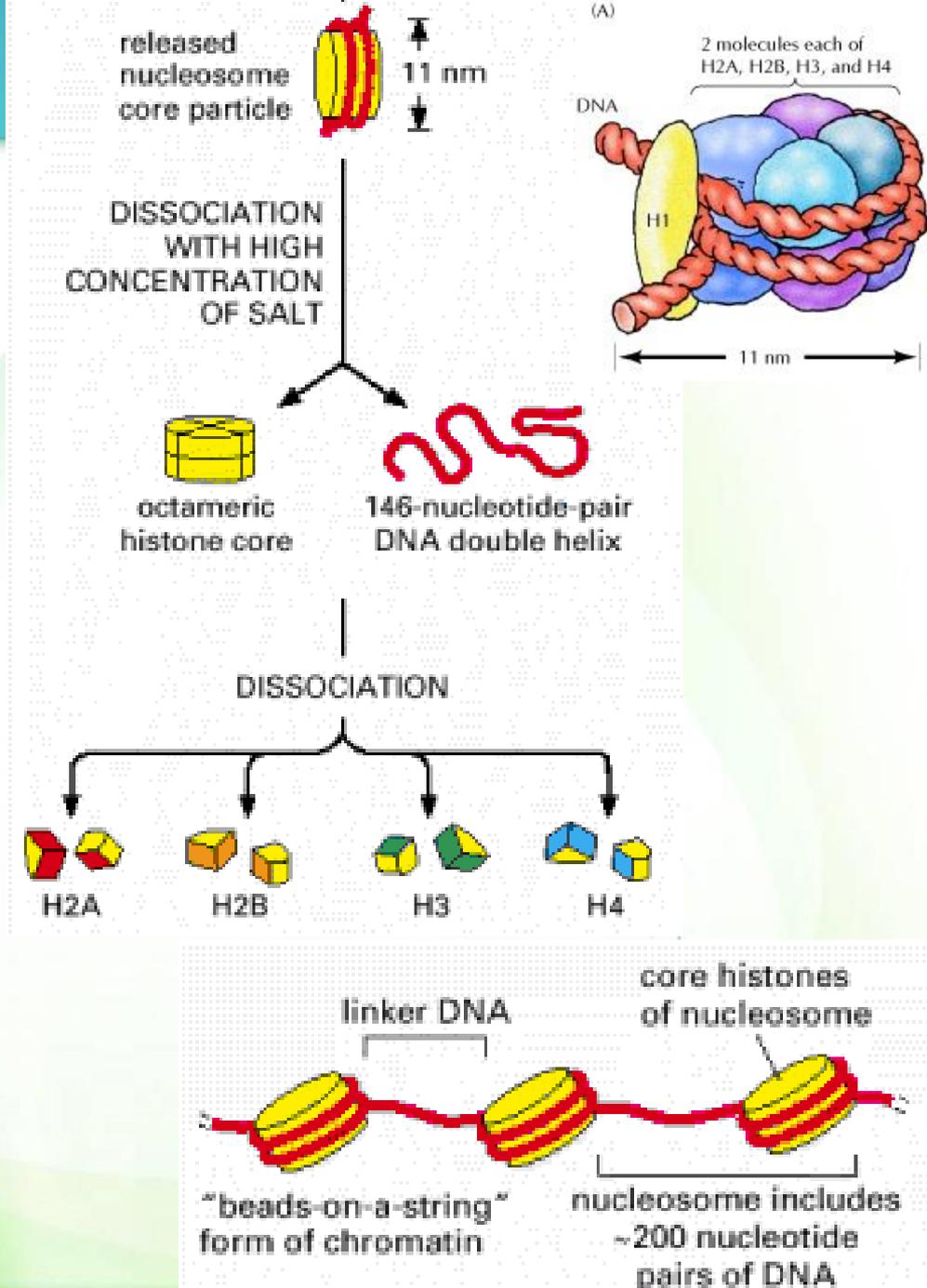


- In eukaryotes, DNA is coiled to package the large DNA and regulate gene activity.
- Eukaryotic DNA is complexed with a number of proteins, principally histones, which are surrounded by DNA.
- Chromatin = DNA molecule + proteins.



Nucleosomes

- The histone protein core is an octamer (two molecules of histones H2A, H2B, H3, and H4).
- A linker DNA/spacer region connects the octamer-DNA complexes.
- A **nucleosome** consists of DNA wrapped around a histone core.
- H1 is bound to the the octamer and wrapped DNA (a **chromatosome**).
- Histones are positively charged facilitating DNA interaction and charge neutralization.



Light absorbance of nucleic acids



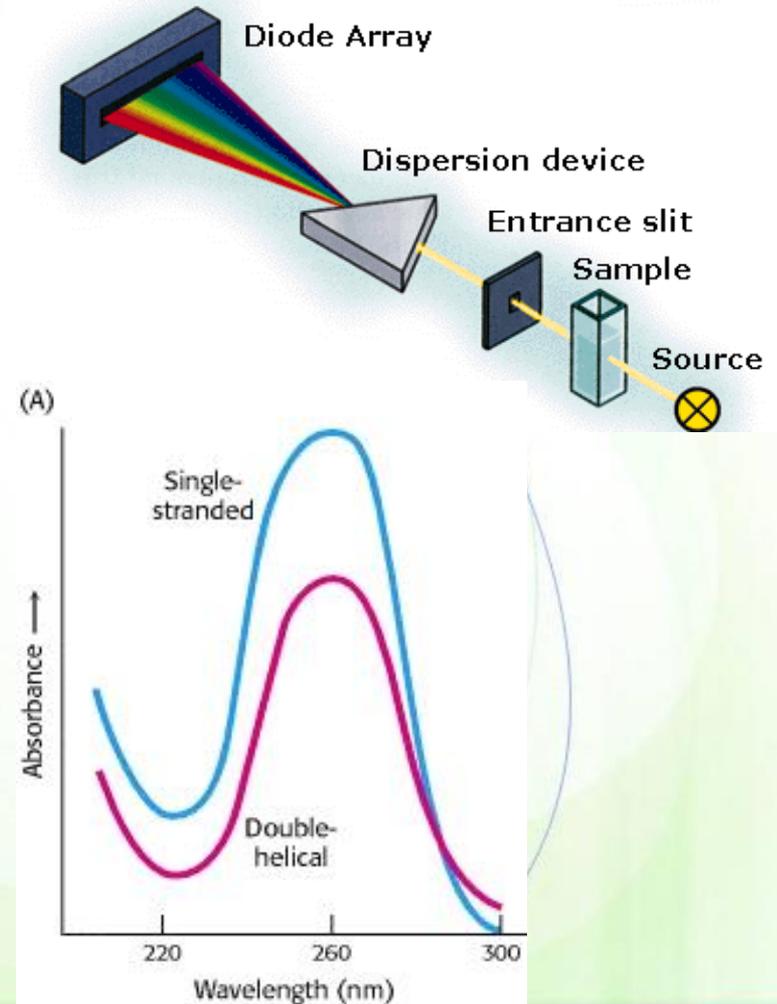
- Aromatic pyrimidines and purines can absorb UV light
- The peak absorbance is at 260 nm wavelength
- The absorbance of nucleic acids at 260 nm (A_{260}) is constant
 - dsDNA: A_{260} of 1.0 = 50 $\mu\text{g/ml}$
 - ssDNA: A_{260} of 1.0 = 30 $\mu\text{g/ml}$
 - ssRNA: A_{260} of 1.0 = 40 $\mu\text{g/ml}$

Reason for ss vs. ds absorbance:

- Unstacked bases vs. stacked bases

What is the concentration of a double stranded DNA sample diluted at 1:10 and the A_{260} is 0.1?

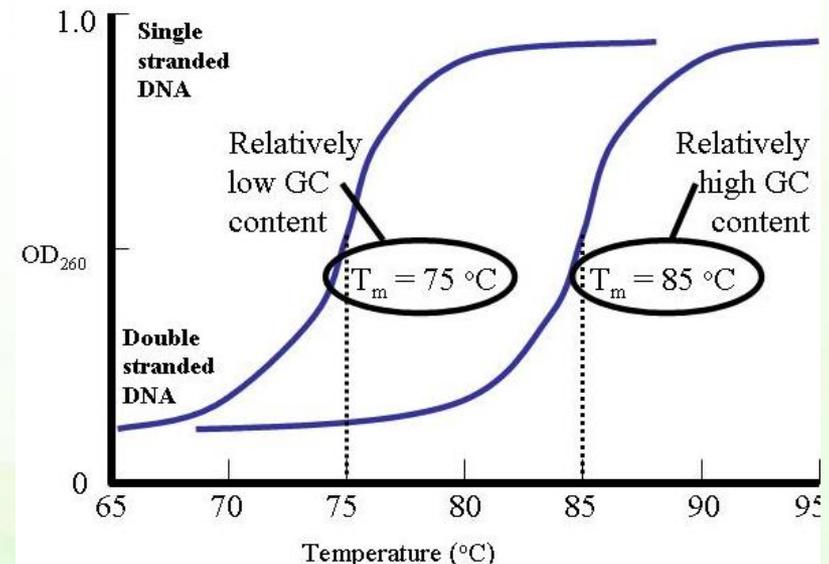
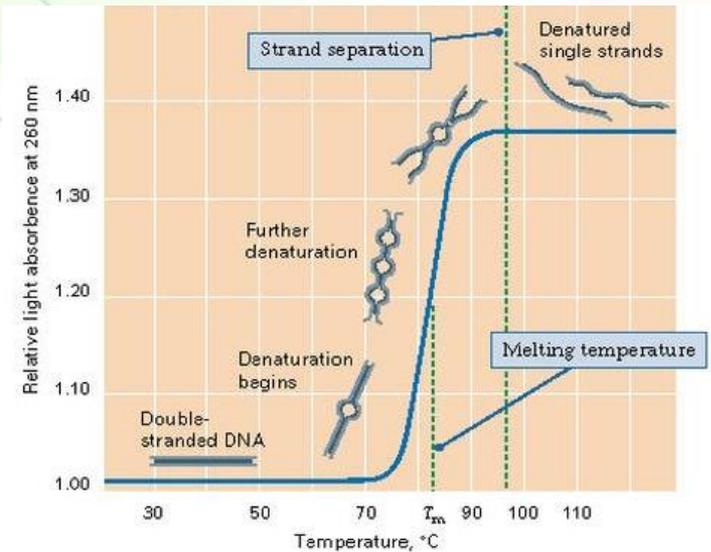
$$\begin{aligned}\text{DNA concentration} &= 0.1 \times 10 \times 50 \mu\text{g/ml} \\ &= 50 \mu\text{g/ml}\end{aligned}$$



Observation of denaturation



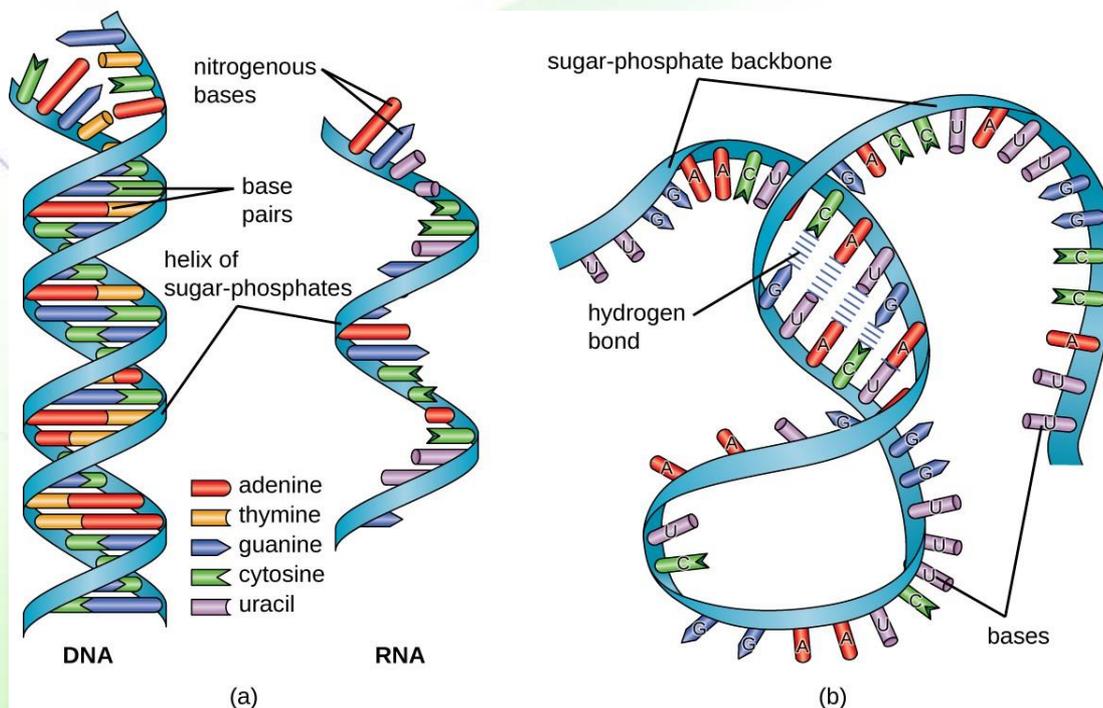
- The transition temperature, or melting temperature (T_m).
- Factors influencing T_m
 - Length
 - G·C pairs
 - Hydrogen bonds
 - Base stacking
 - pH
 - Salts and ions
 - Destabilizing agents (alkaline solutions, formamide, urea)



RNA



- It consists of long, unbranched chains of nucleotides joined by phosphodiester bonds between the 3'-OH of one pentose and the 5'-PO₄⁻ of the next.
- The pentose unit is β-D-ribose (it is 2-deoxy-D-ribose in DNA).
- The pyrimidine bases are uracil and cytosine (thymine and cytosine in DNA).
- In general, RNA is single stranded (DNA is double stranded).



RNA does not have a precise structure, but it can fold on itself forming hydrogen bonds within the same molecule.

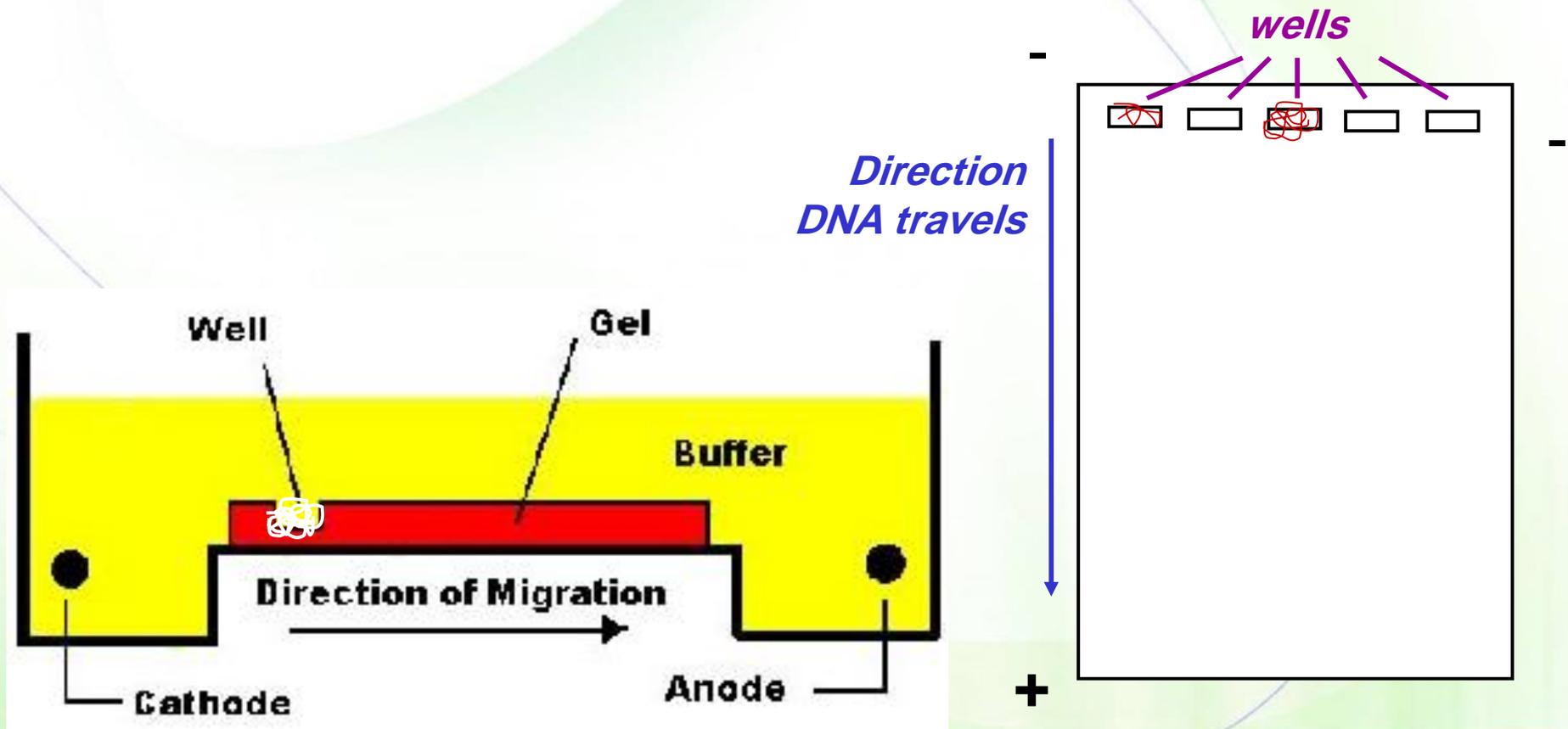
Types of RNA



Non-coding RNA	Length (nt)	Species	Function
Ribosomal RNA (rRNA)	120~4700	All	Translation
Transfer RNA (tRNA)	70~100	All	Translation
Small nuclear RNA (snRNA)	70~350	Eukaryote	Splicing, mRNA processing
Small nucleolar RNA (snoRNA)	70~300	Eukaryote, archaea	RNA modification, rRNA processing
miRNA	21~25	Eukaryote	Translational regulation
siRNA	21~25	Eukaryote	Protection against viral infection
piRNA	24~30	Eukaryote	Genome stabilization
Long ncRNA	several hundreds~ several hundred thousands	Eukaryote	Transcription, splicing, transport regulation

Gel electrophoresis

- The length and purity of DNA molecules can be accurately determined by the gel electrophoresis.



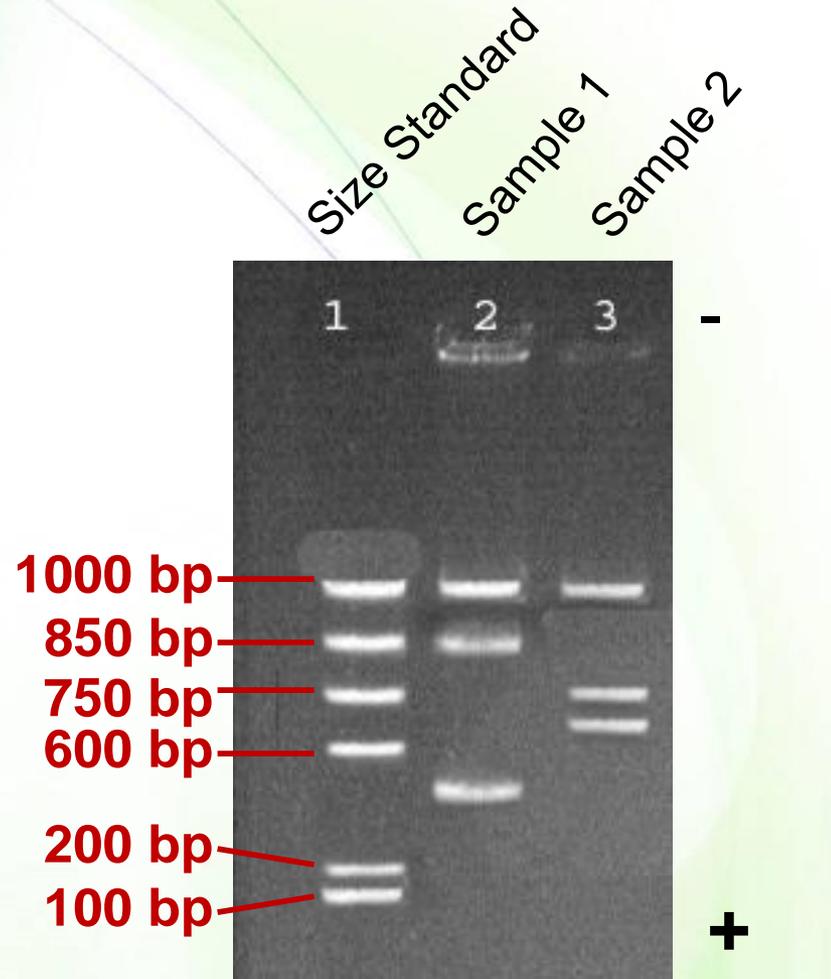
Resources



- <http://www.personal.psu.edu/pzb4/electrophoresis.swf>
- <http://www.sumanasinc.com/webcontent/animations/content/gelectrophoresis.html>
- <http://www.sumanasinc.com/webcontent/animations/content/gelectrophoresis.html>

Detection

- The DNA molecules of different lengths will run as "bands".
- Each bands contains thousands to millions of copies of DNA fragments of the same length. They can be of same or different type (not one DNA molecule).
- DNA is **stained** (that is, colored) with a dye (ethidium bromide) or radioactively **labeled** (^{32}P).
- It is common that a DNA standard is used to determine the length of the examined DNA molecule.

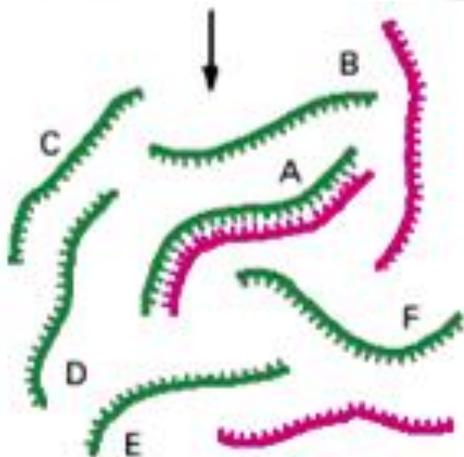
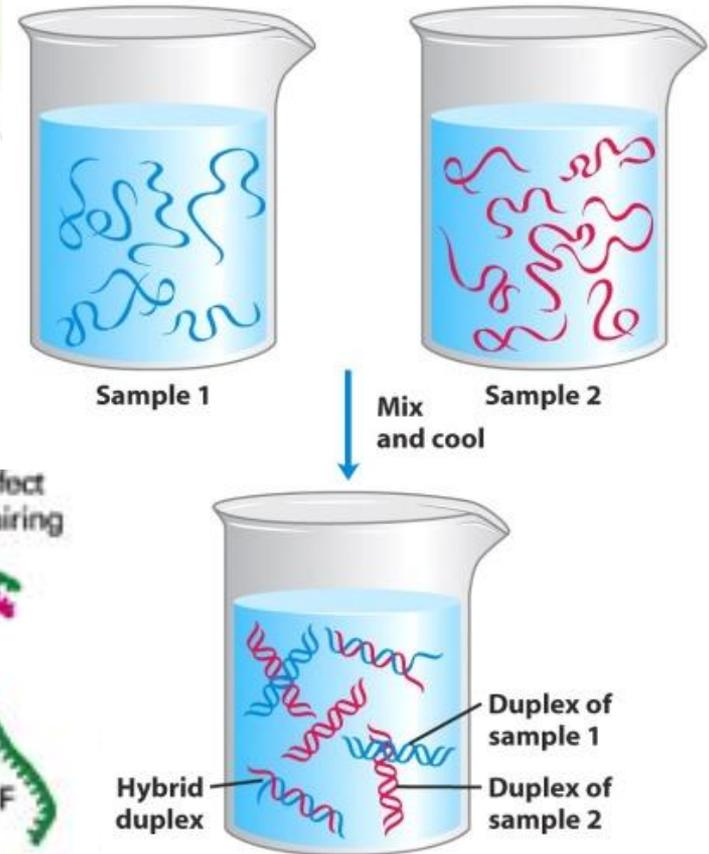


bp: base pair

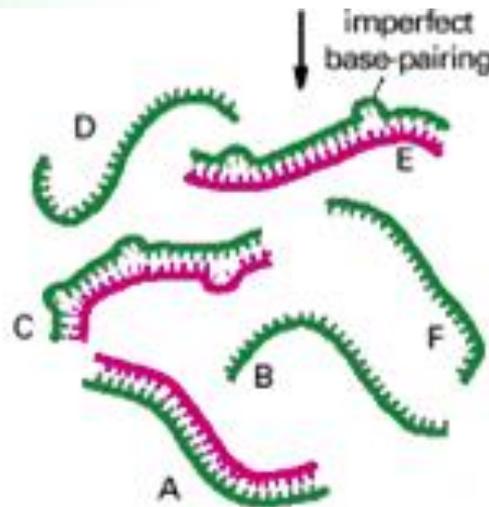
Hybridization



- DNA from different sources can form double helix as long as their sequences are compatible (hybrid DNA).
- Hybridization can be imperfect.



only A forms stable double helix



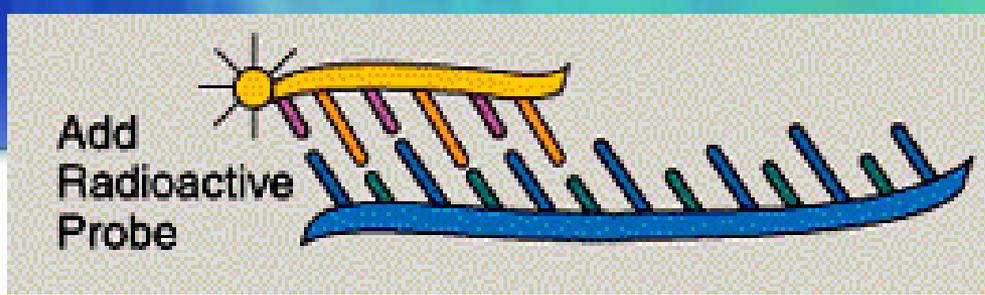
A, C, and E all form stable double helices

Hybridization techniques



- Hybridization reactions can occur between any two single-stranded nucleic acid chains provided that they have complementary nucleotide sequences
- Hybridization reactions are used to detect and characterize specific nucleotide sequences

Probes



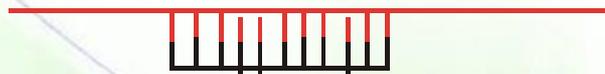
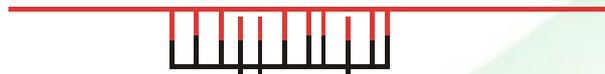
- A probe is a short sequence of single stranded DNA (an oligonucleotide) that is complementary to a small part of a larger DNA sequence.
- Hybridization reactions use labeled DNA probes to detect larger DNA fragments.



Hybridization can be (non)specific



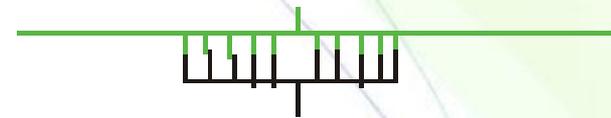
Perfectly
Complementary



Duplex

Conditions
near T_m

Mismatch



+

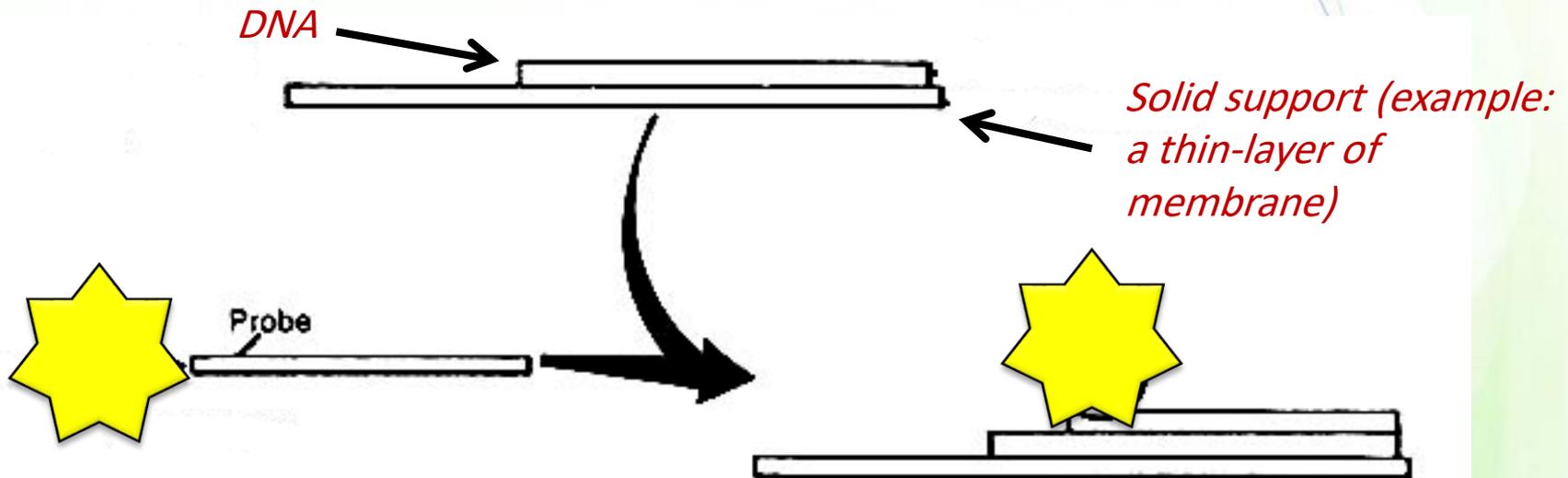


No duplex

Dot blot



- This is a technique that informs us if a specific sequence that is complementary to a probe of a known sequence exists in a larger DNA.
- DNA is bound to a solid support and a labeled probe is added. If binding occurs, the sequence exists.



Disease detection by ASO (Cystic fibrosis)



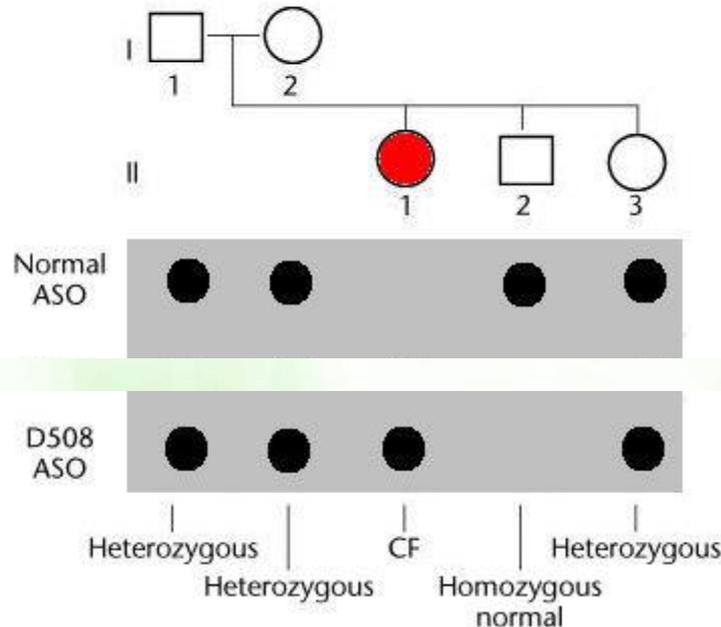
ASO: Allele-specific oligonucleotide

The whole genomic DNA is spotted on a solid support (like a nylon membrane) and hybridized with two ASO's, one at a time.

Cystic Fibrosis allele $\Delta 508$ has 3bp deletion [AGA]

ASO for normal DNA 5' CACCAA[AGA]GATATTTTC-3'

ASO for DNA sequence of $\Delta 508$ mutation 5' CACCAATGATATTTTC-3'



Southern blotting



- This technique is a combination of DNA gel electrophoresis and hybridization
- Used to detect:
 - the presence of a DNA segment complementary to the probe
 - the size of the DNA fragment

