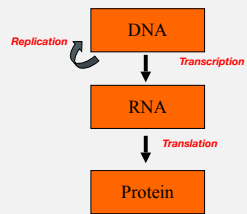


BIOMOLECULES

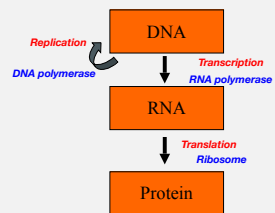
ECS129

Instructor: Patrice Koehl

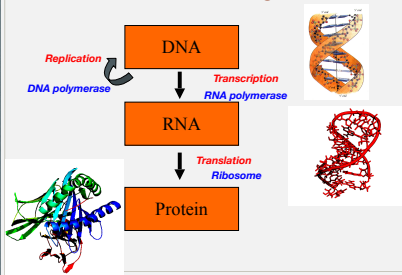
Central Dogma



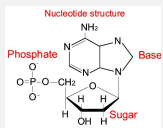
Central Dogma



Central Dogma



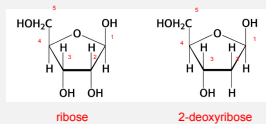
DNA



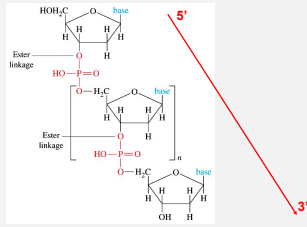
Base Formula	Base (XHT)	Nucleotide Ribose or deoxyribose	Nucleotide X-ribose phosphate X=base phosphate
<chem>NC1=NC=CC(=N1)</chem>	Cytosine, C	Cytosine, A	Cytosine nucleoside diphosphate CDP
<chem>C1=CN(C=C1)C(=O)N</chem>	Thymine, T	Thymine, T	Thymine nucleoside diphosphate TDP
<chem>NC1=NC=NC2=C1N=CN2</chem>	Adenine, A	Adenosine, A	Adenosine nucleoside diphosphate ADP
<chem>C1=NC2=C(N1)C(=O)N(C)N2</chem>	Guanine, G	Guanosine, A	Guanosine nucleoside diphosphate GDP

DNA

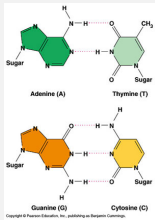
These "bases" are attached to sugar rings: ribose (RNA), deoxyribose (DNA):



DNA



DNA



In other words, if an adenine forms one member of a pair, on either chain, then on the complementary the other member must be thymine - similarly for guanine and cytosine. The sequence of bases on a single chain does not appear to be restricted in any way. However, if only specific pairs of bases can be formed, it follows that if the sequence of bases on one chain is given, then the sequence on the other chain is automatically determined.

It has not escaped our notice that the specific pairing we have postulated immediately suggests a possible copying mechanism for the genetic material.

Excerpt from Watson and Crick, Nature, 436, 737-728 (1953)

DNA

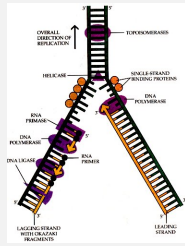


We wish to put forward a radically different structure for the salt of deoxyribose nucleic acid. This structure has two helical chains each coiled round the same axis (see diagram). We have made the usual chemical assumptions, namely, that each chain consists of phosphate di-ester groups joining β -D-deoxy-ribofuranose residues with 3',5' linkages. The two chains (but

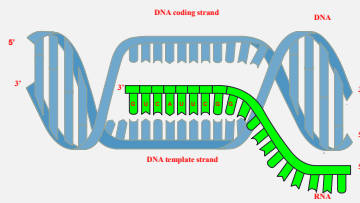
Excerpt from Watson and Crick, Nature, 436, 737-728 (1953)

DNA → DNA

- **Helicase** separates the two DNA strands, starting at replication origins (rich in A-T base pairs)
- **RNA primase** inserts a starter of RNA nucleotides at the initiation point
- **DNA polymerase** binds a complementary leading strand of DNA nucleotides starting at the 3' end of the RNA prime

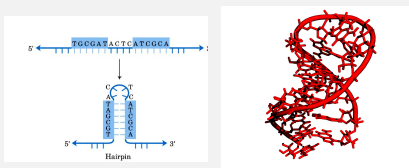


DNA → RNA



RNA

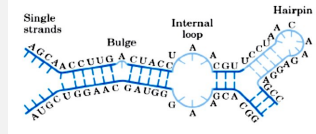
Single stranded subsequences bounded by base pairs are called **loops**. A loop at the end of a stem is called a **hairpin loop**. Simple substructures consisting of a single stem and loop are called **stem loops**, or **hairpins**.



RNA

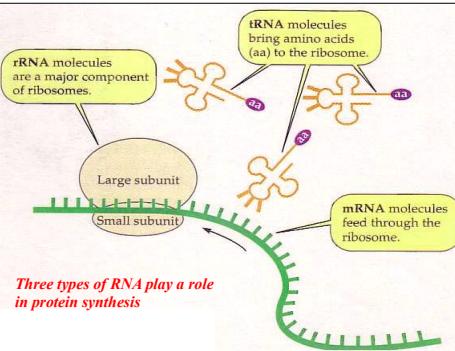
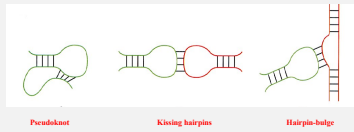
Single stranded bases within a stem form a **bulge** or **bulge loop** if the single stranded bases are on only one side of the stem.

If single stranded bases interrupt both sides of a stem, they form an **internal (interior) loop**.



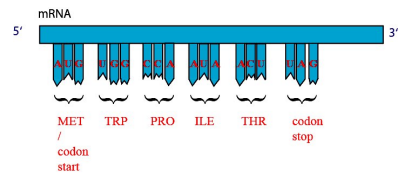
RNA

In addition to secondary structural interactions in RNA, there are also tertiary interactions, including: (A) **pseudoknots**, (B) **kissing hairpins** and (C) **hairpin-bulge** contacts.



TRANSLATION

- The process of reading the mRNA sequence and creating the protein is called **translation**
- Proteins are made of amino acids (20 different, 9 "essentials")
- 3 bases or nucleotides make one **codon**
- Each codon specifies one amino acid - genetic code

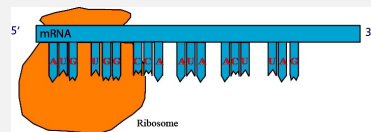


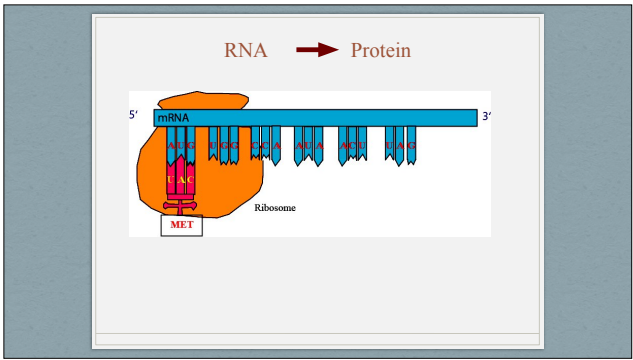
RNA → Protein

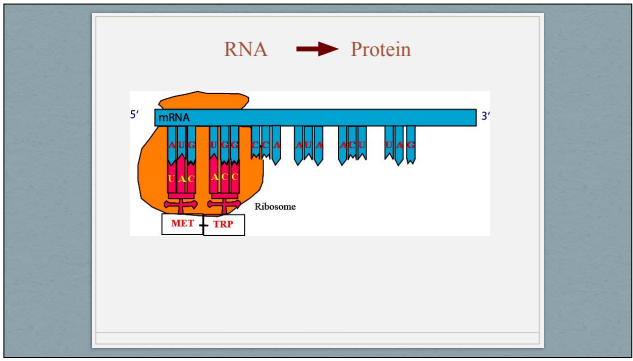
		2 nd base in codon				
		U	C	A	G	
1 st base in codon	U	Phe Phe Leu Leu	Ser Ser Ser STOP	Tyr Tyr STOP Tip	Cys Cys STOP Tip	U C A G
	C	Leu Leu Leu	Pro Pro Pro	His His Gln	Arg Arg Arg	U C A G
	A	Ile Ile Ile Met/Start	Thr Thr Thr	Asn Asn Lys	Ser Ser Arg Arg	U C A G
G	Val Val Val Val	Ala Ala Ala Ala	Asp Asp Glu Glu	Gly Gly Gly Gly	U C A G	

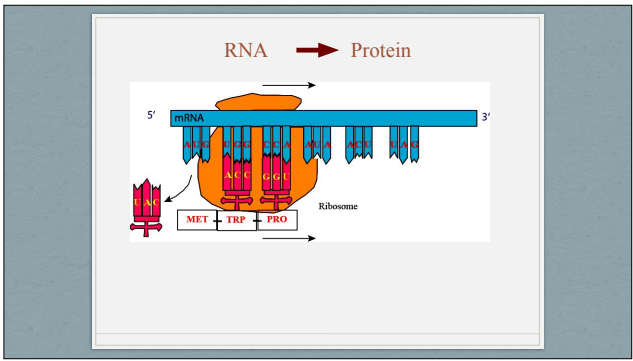
3rd base in codon

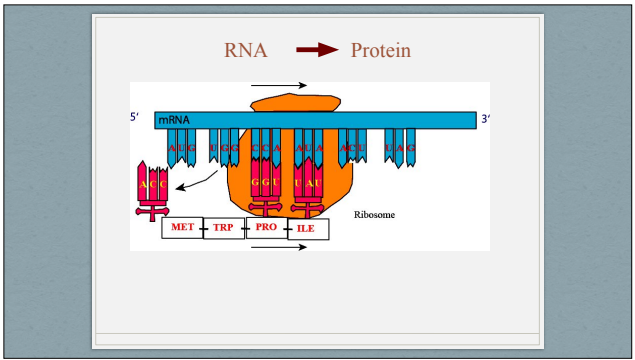
RNA → Protein

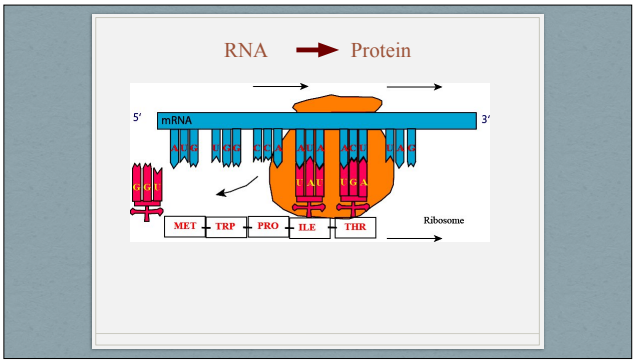


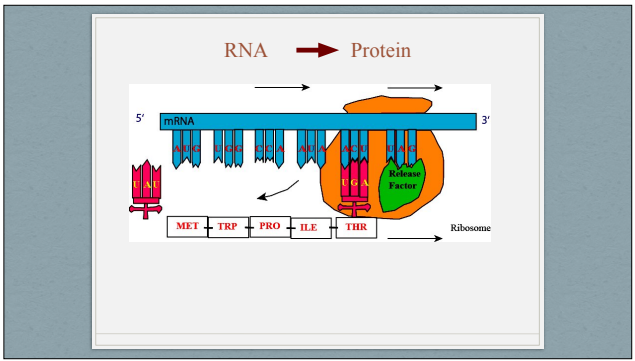


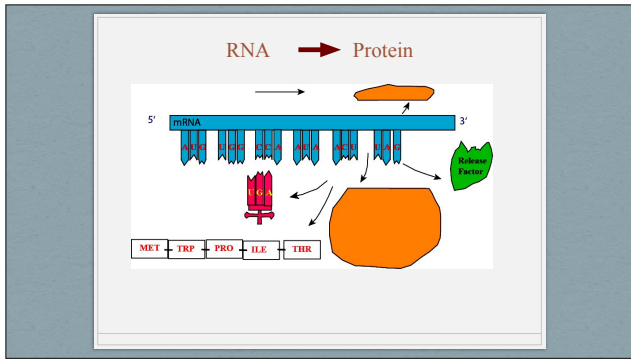


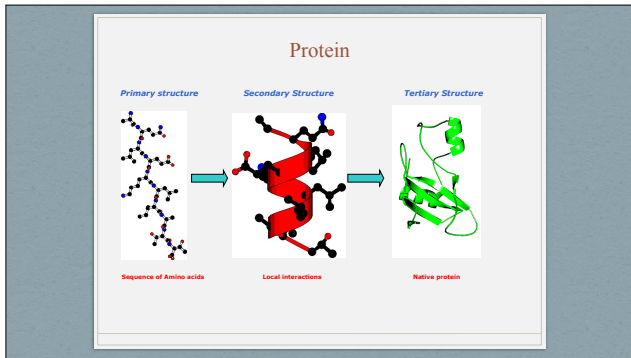


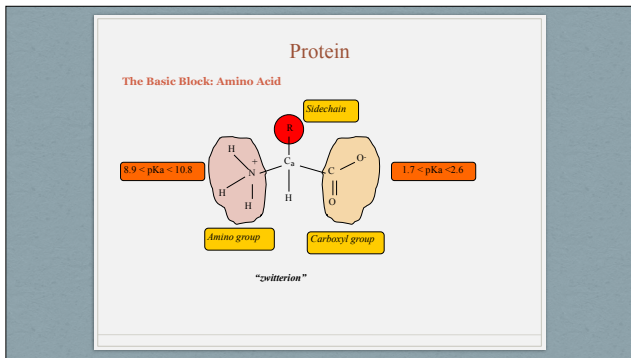












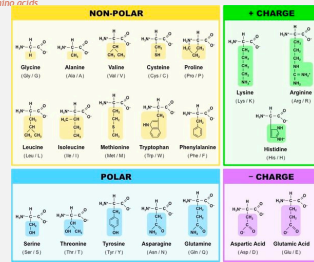
Protein

The 20 amino acids

1-letter	3-letter	Amino acid
A	Ala	Alanine
C	Cys	Cysteine
D	Asp	Aspartic Acid
E	Glu	Glutamic Acid
F	Phe	Phenylalanine
G	Gly	Glycine
H	His	Histidine
I	Ile	Isoleucine
K	Lys	Lysine
L	Leu	Leucine
M	Met	Methionine
N	Asn	Asparagine
P	Pro	Proline
Q	Gln	Glutamine
R	Arg	Arginine
S	Ser	Serine
T	Thr	Threonine
V	Val	Valine
W	Trp	Tryptophane
Y	Tyr	Tyrosine

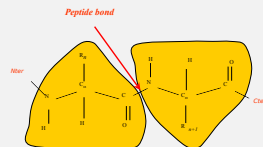
Protein

The 20 amino acids



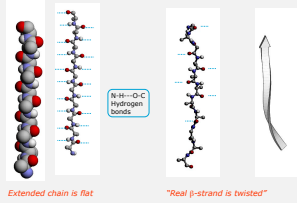
Protein

The Protein: A polymer of Amino acids



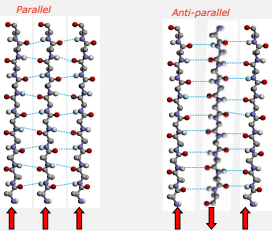
Protein

The β -strand



Protein

Two types of β -sheets



Protein

Protein Tertiary Structure

- All α proteins
- All β proteins
- Alpha and beta proteins:
 - α/β proteins (alternating α and β)
 - $\alpha \leftrightarrow \beta$ proteins

