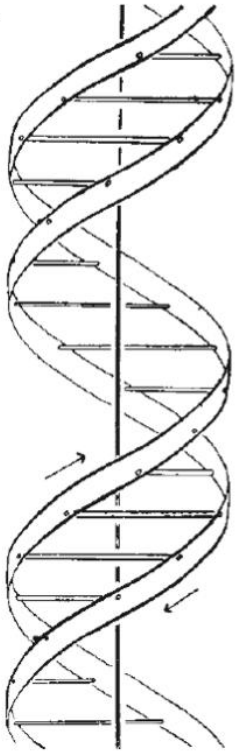
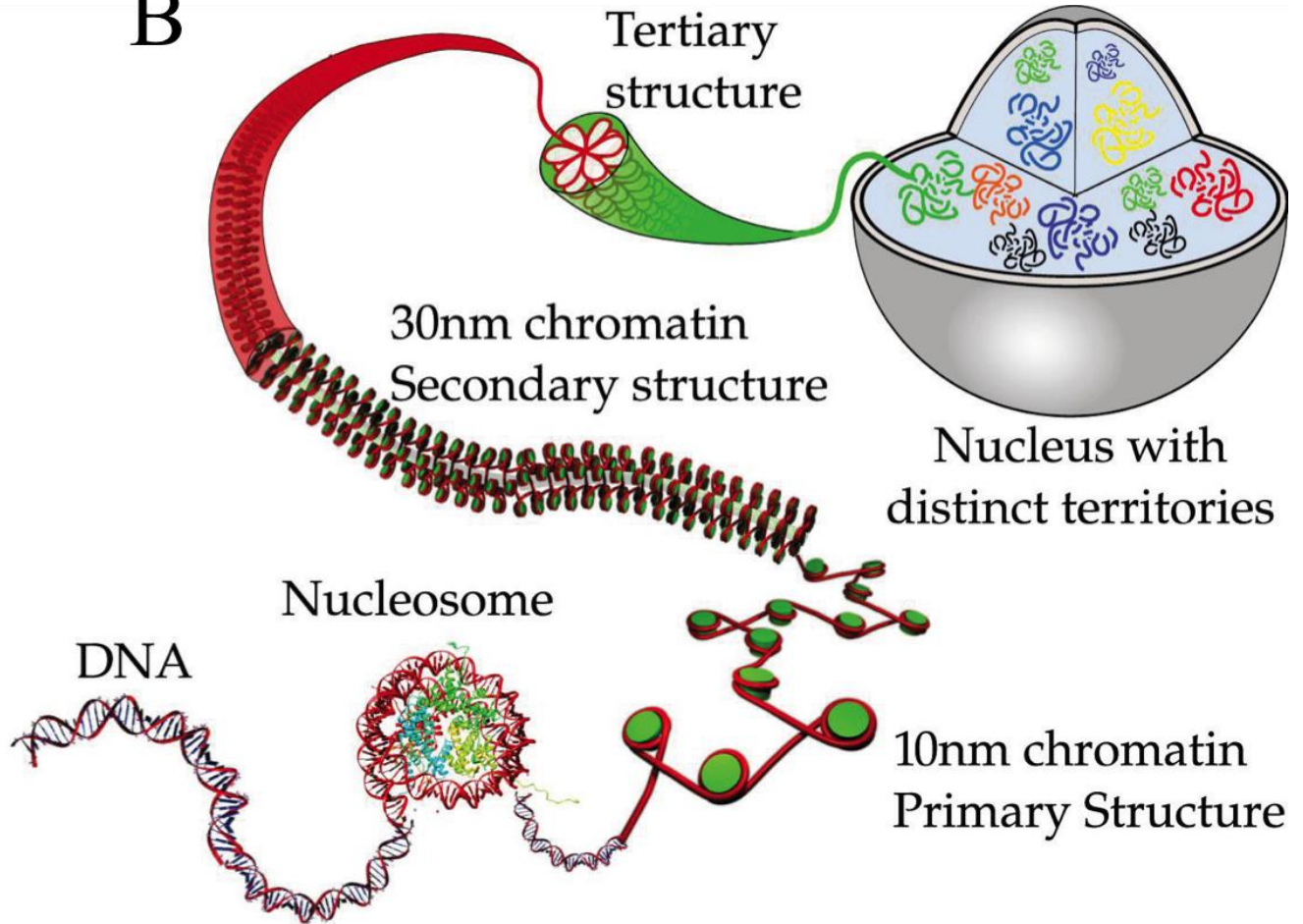


A



This figure is purely diagrammatic. The two ribbons symbolize the two phosphate—sugar chains, and the horizontal rods the pairs of bases holding the chains together. The vertical line marks the fibre axis

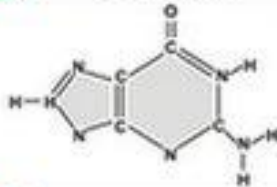
B



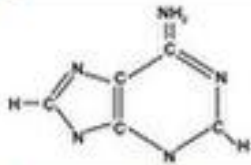
**Cytosine**



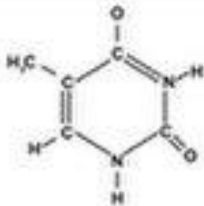
**Guanine**



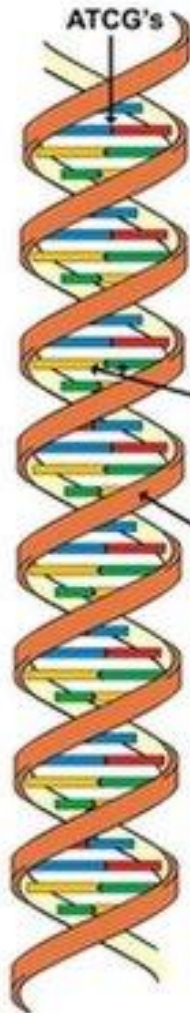
**Adenine**



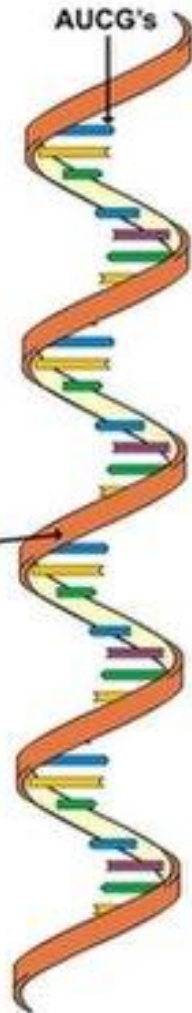
**Thymine**



Nitrogenous Bases

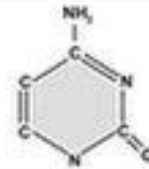


**DNA**  
Deoxyribonucleic Acid



**RNA**  
Ribonucleic Acid

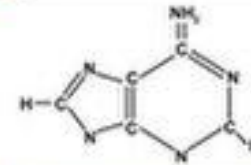
**Cytosine**



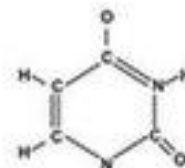
**Guanine**



**Adenine**



**Uracil**



Replaces Thymine in RNA

Nitrogenous Bases

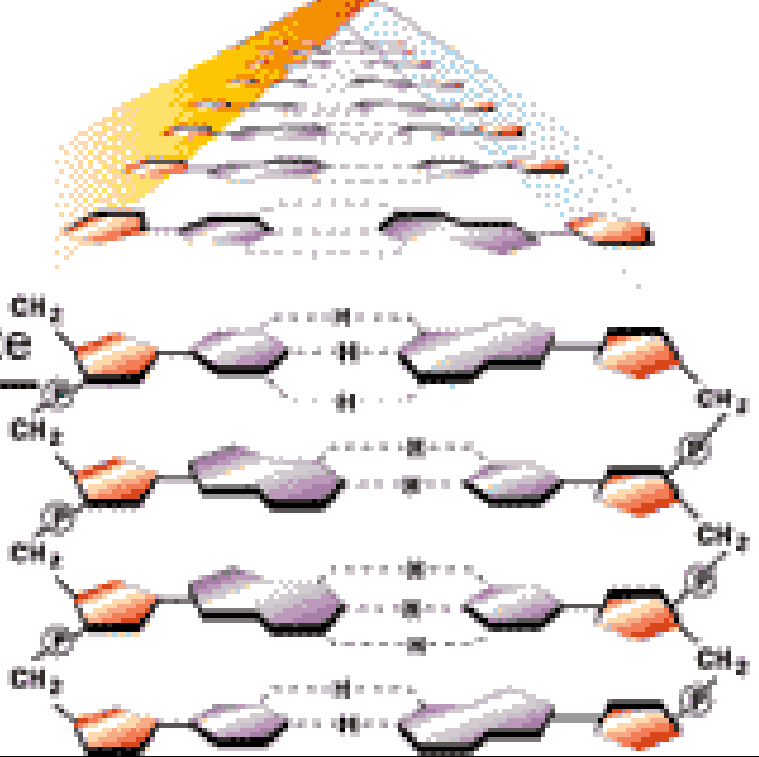
# DNA Double Helix



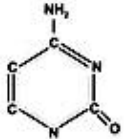
Sugar — 

Bases — 

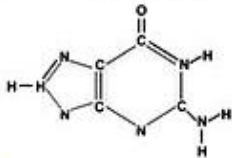
Phosphate  
Group



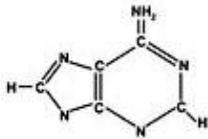
● **Cytosine**



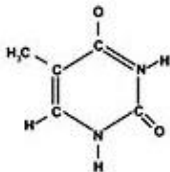
● **Guanine**



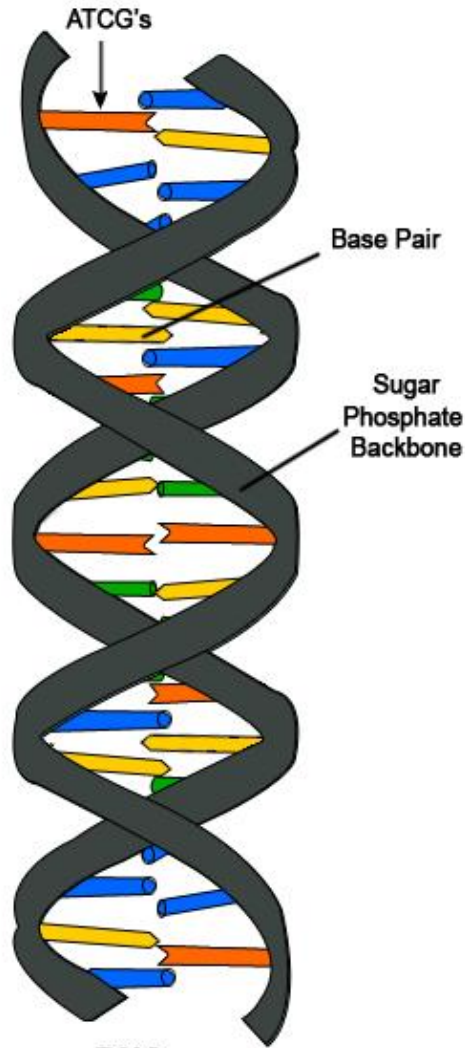
● **Adenine**



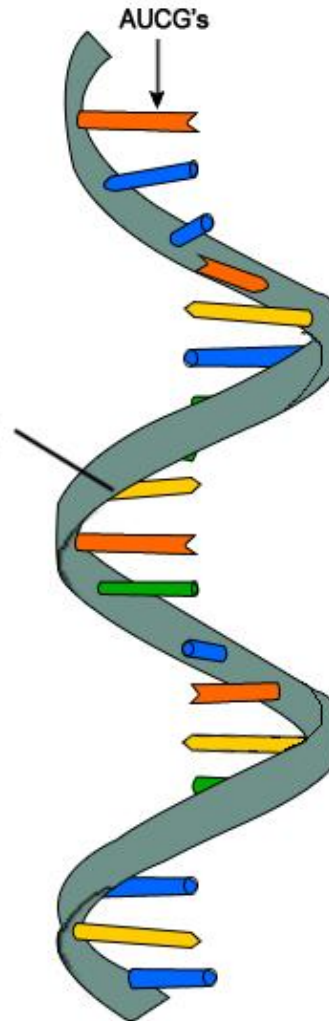
● **Thymine**



Nitrogenous  
Bases

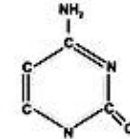


**DNA**  
Deoxyribonucleic Acid

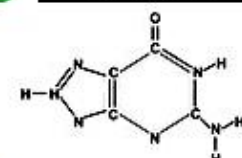


**RNA**  
Ribonucleic Acid

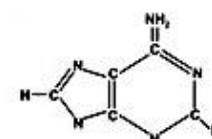
● **Cytosine**



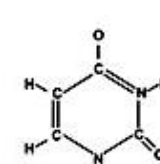
● **Guanine**



● **Adenine**



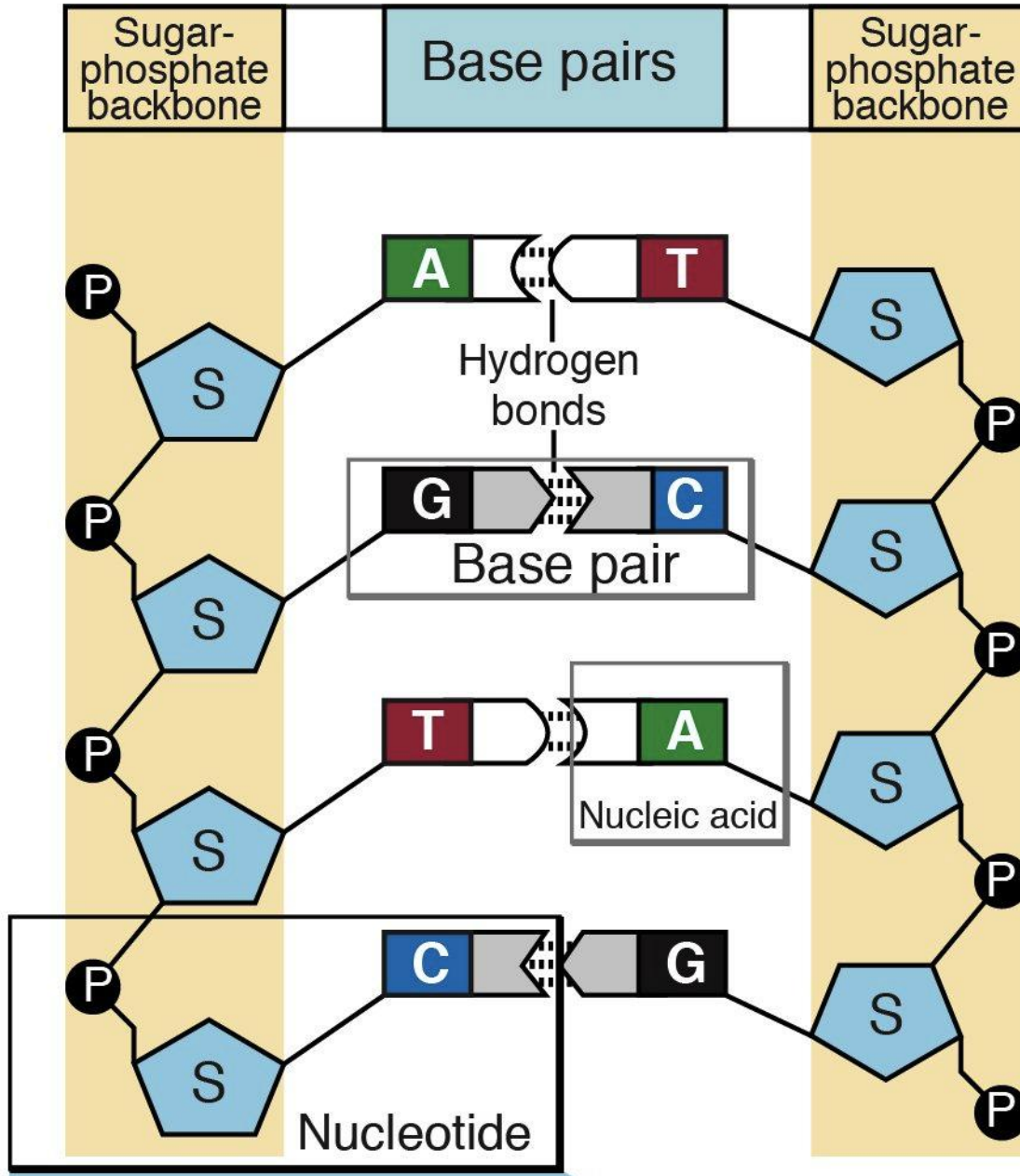
● **Uracil**



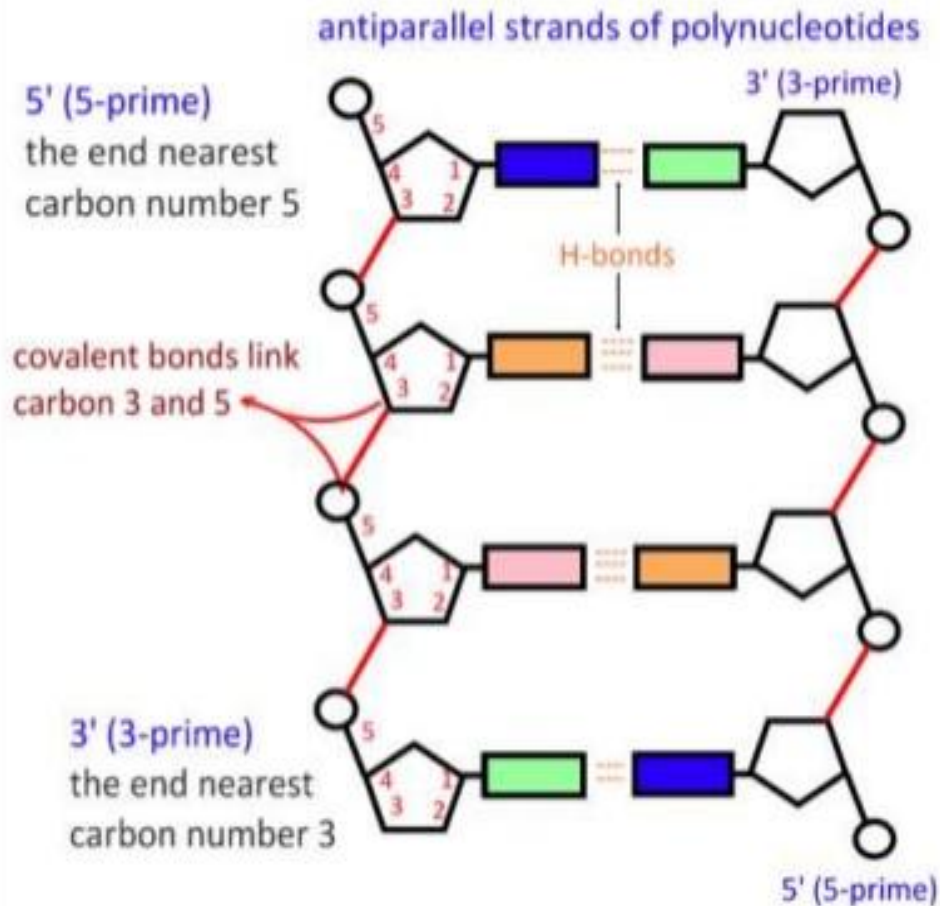
Replaces Thymine in RNA

Nitrogenous  
Bases

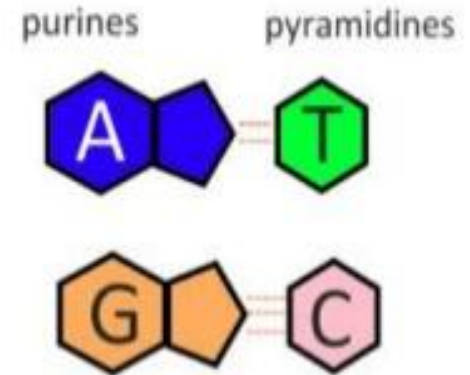
# Deoxyribonucleic Acid (DNA)



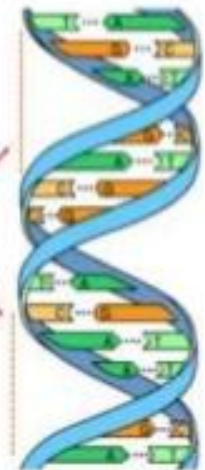
**Core Review:** 2.6.U3 DNA is a double helix made of two antiparallel strands of nucleotides linked by hydrogen bonding between complementary base pairs.



A purine must base-pair  
with a pyrimidine:

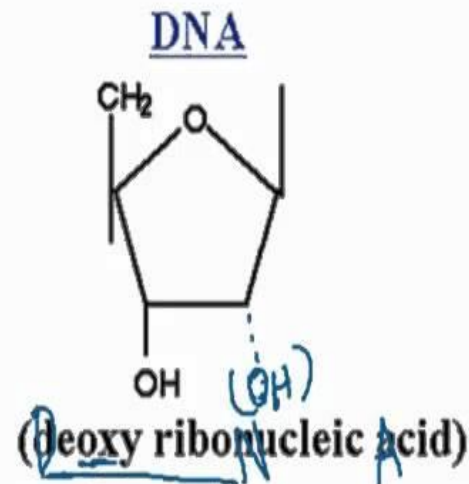
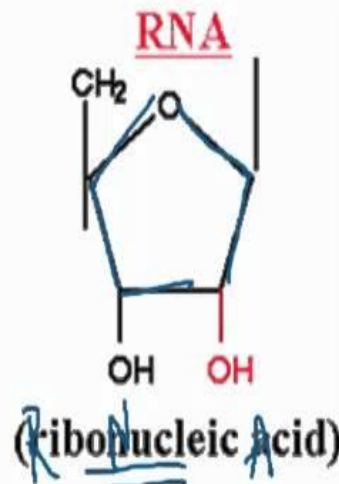


Hydrogen bonds  
also hold the  
structure of the  
double helix

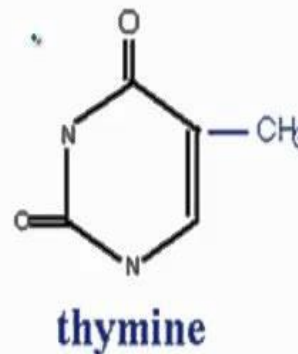
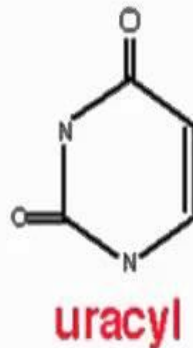


# What makes DNA and RNA different?

1) ribose sugar



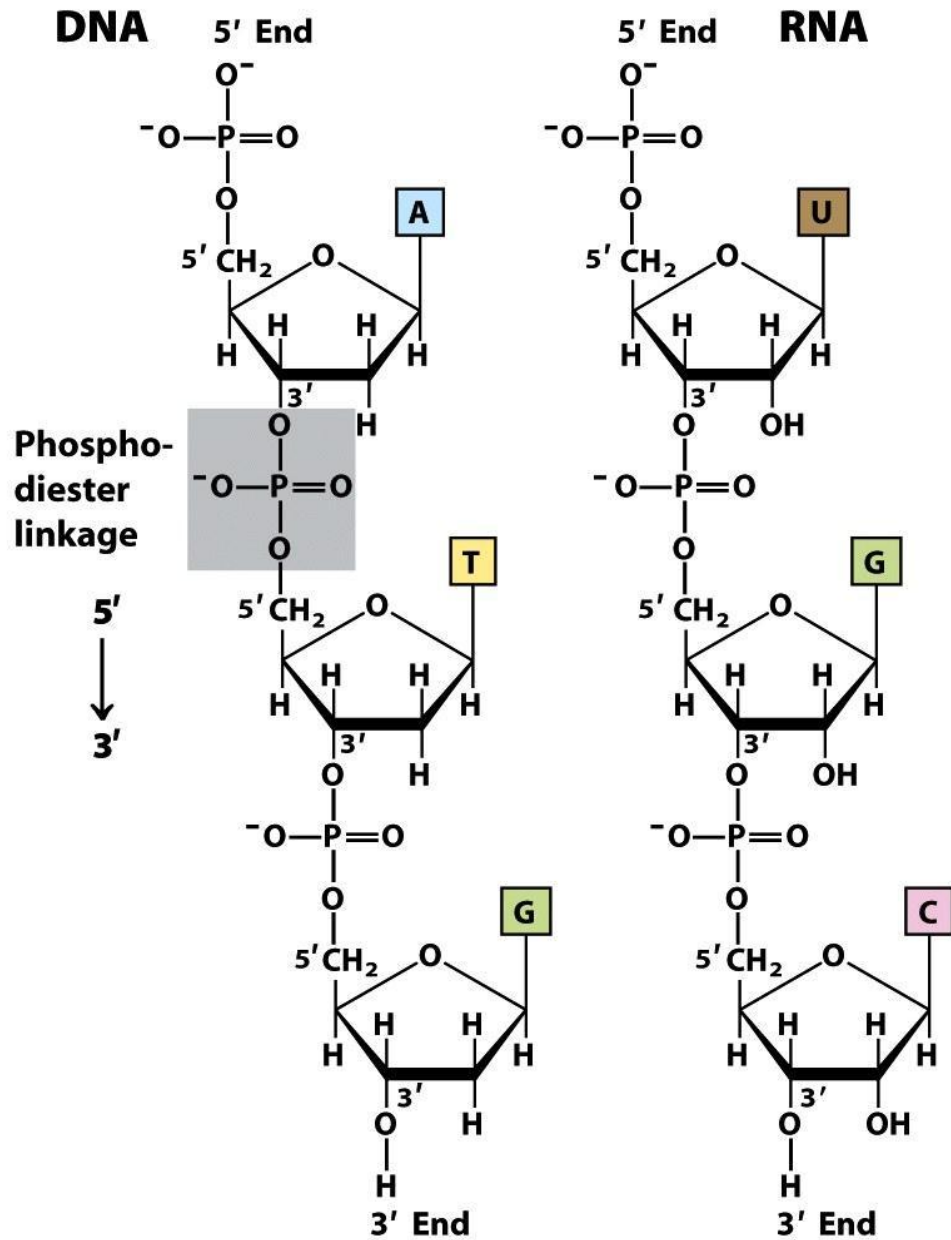
2) T and U



3) strand

single

double

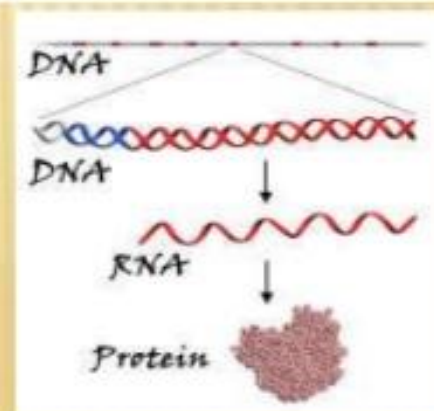
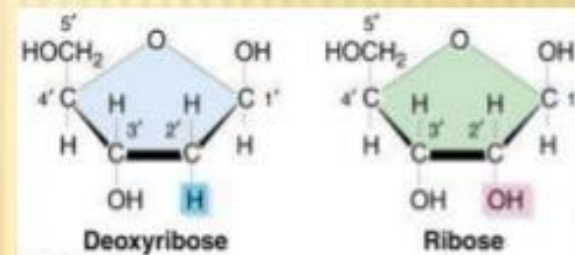
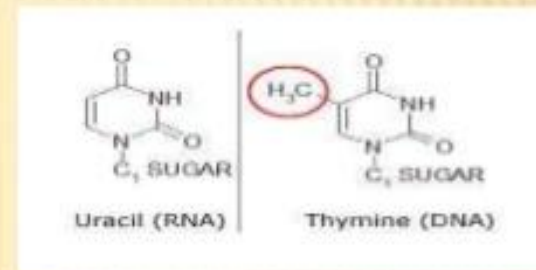


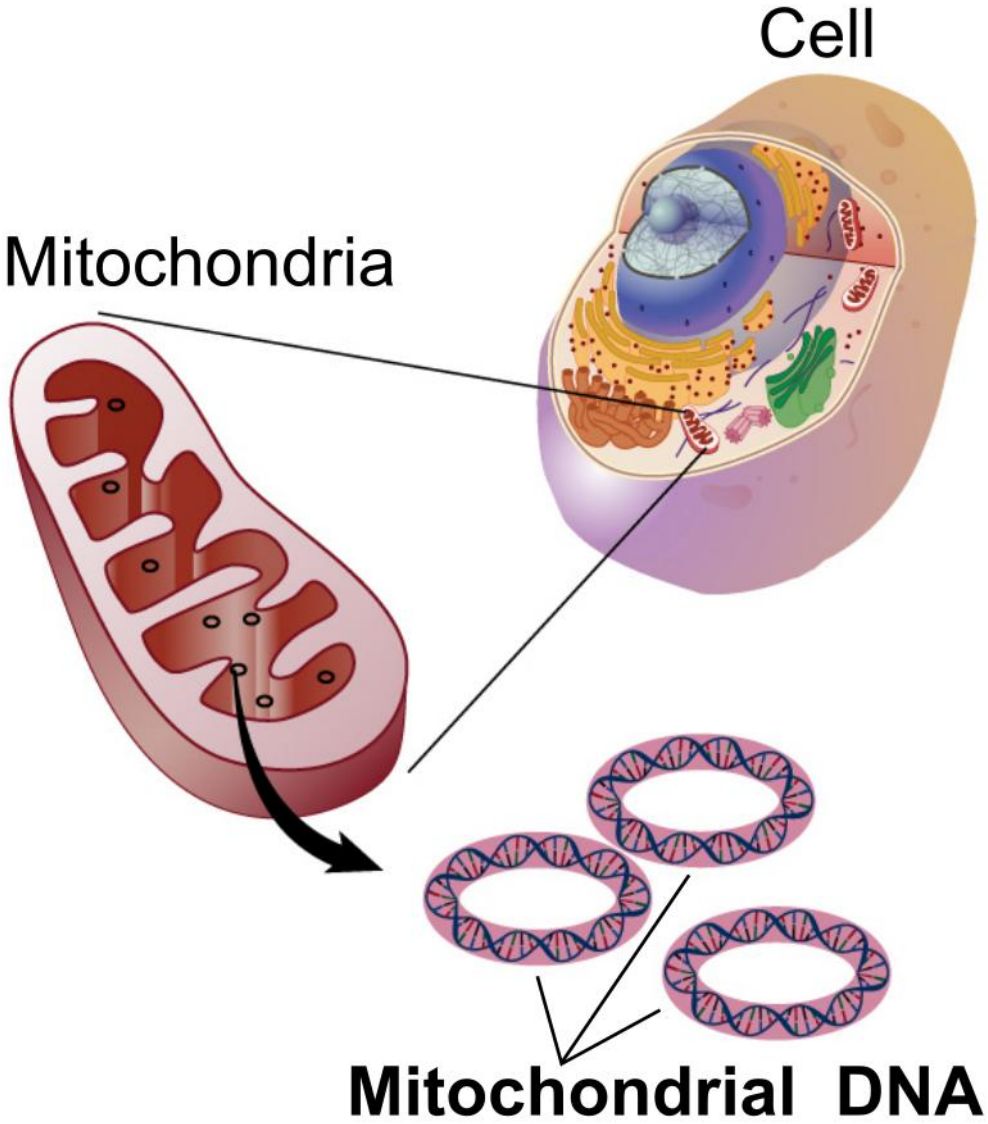
**Figure 8-7**  
*Lehninger Principles of Biochemistry, Fifth Edition*  
 © 2008 W. H. Freeman and Company

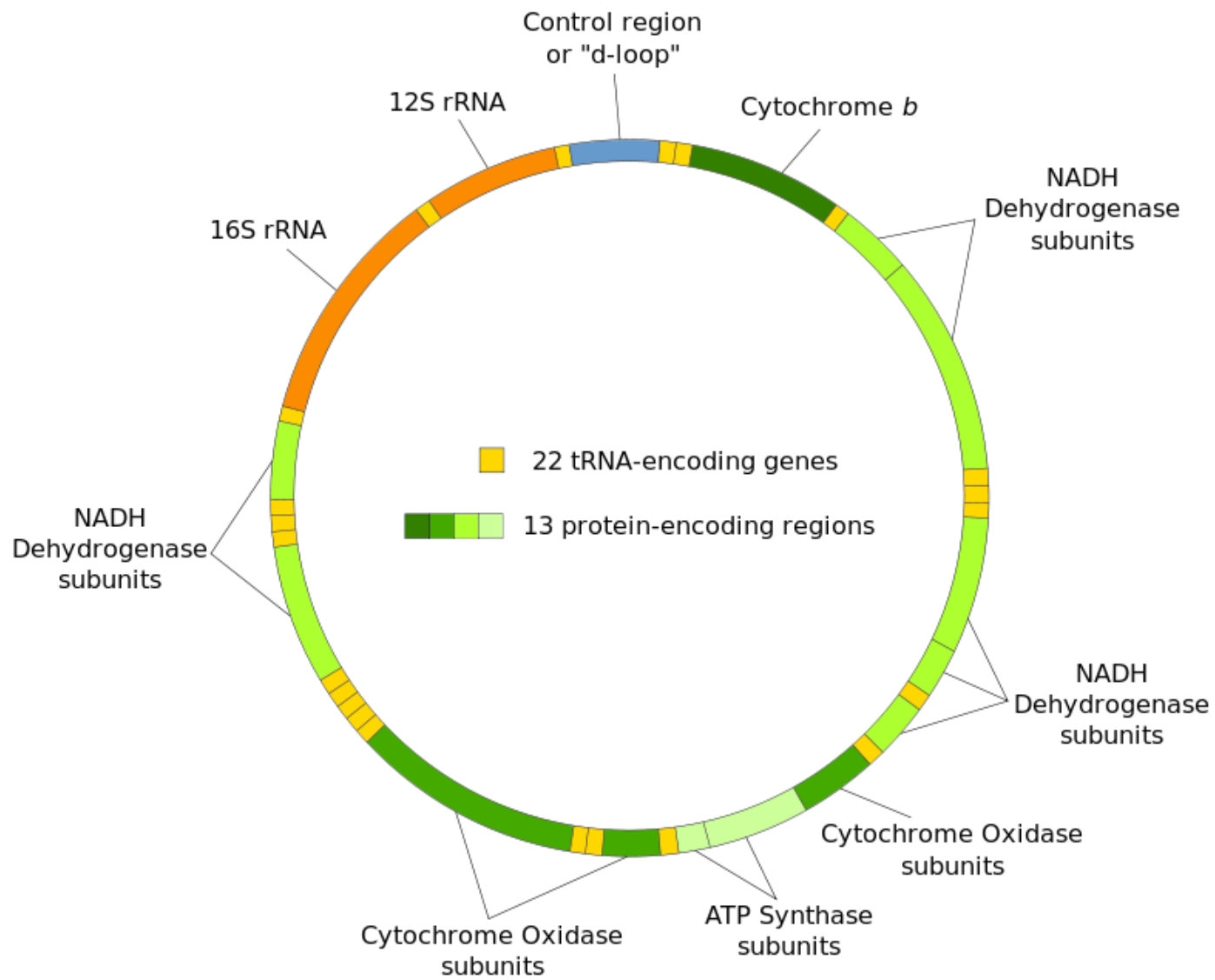


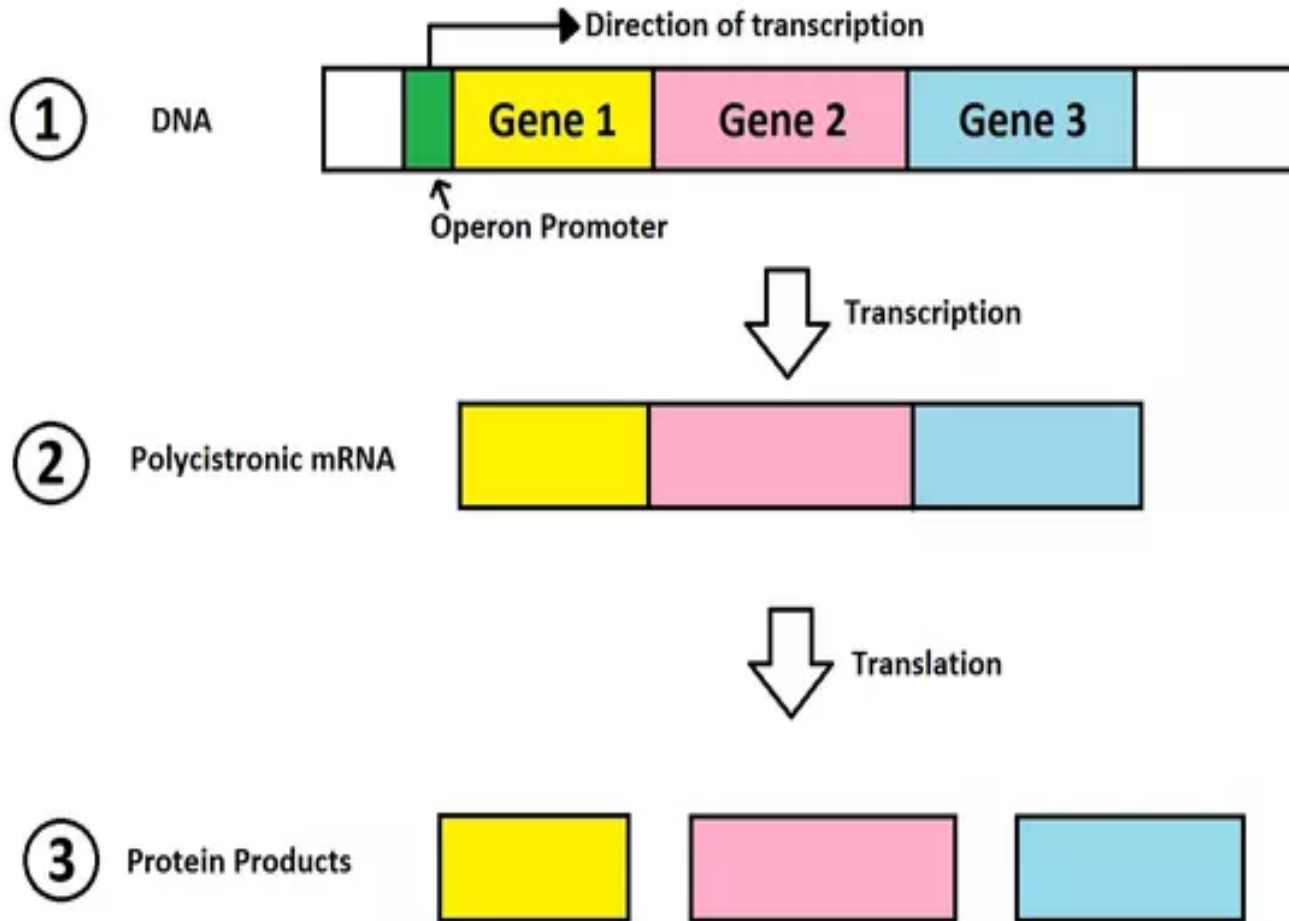
# COMPARATIVE ACCOUNT :DNA AND RNA

	DNA	RNA
Function	Medium of long-term <b>storage</b> and transmission of genetic information	<b>Transfer</b> the genetic code needed for the creation of proteins from the nucleus to the ribosome
Predominant structure	<b>Double- stranded</b> molecule	A <b>single-stranded</b> molecule
Bases	A, T, C, G	A, U, C, G
Sugar	<b>2'- deoxyribose</b>	<b>ribose</b>
Stability	Deoxyribose sugar in DNA is less reactive because of C-H bonds. <b>Stable in alkaline conditions.</b> DNA has smaller grooves, which makes it harder for enzymes to "attack" DNA.	Ribose sugar is more reactive because of C-OH (hydroxyl) bonds. <b>Not stable in alkaline conditions.</b> RNA has larger grooves, which makes it easier to be attacked by enzymes
Propagation	DNA is self-replicating.	RNA is synthesized from DNA when needed

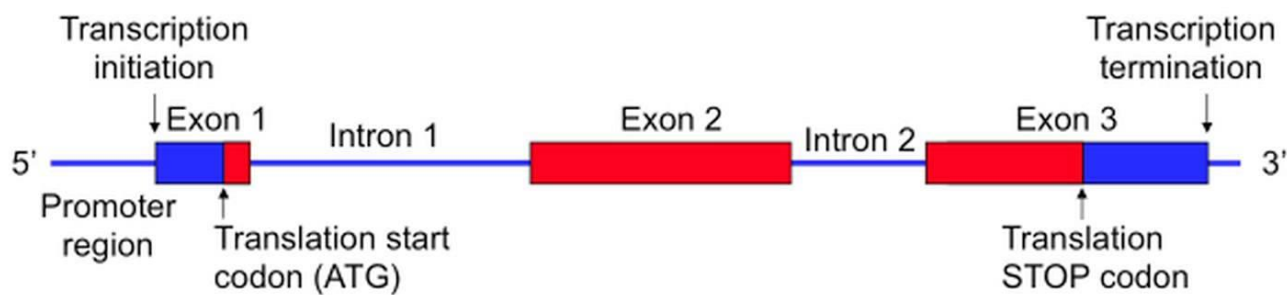


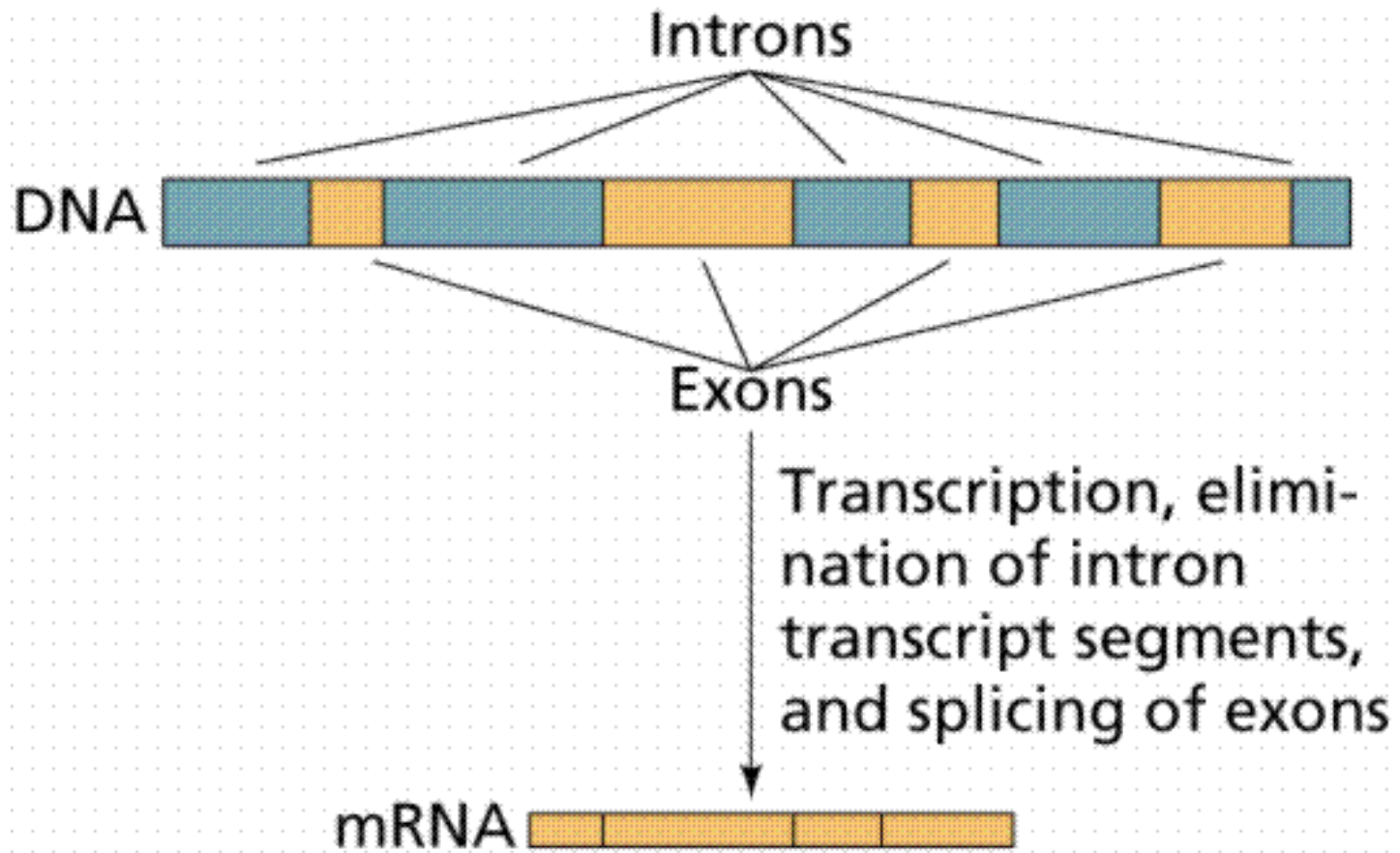






# Gene Structure





## **What is Exons and Introns?**

Exons and Introns: In most eukaryotic genes, coding regions (exons) are interrupted by noncoding regions (introns). During transcription, the entire gene is copied into a pre-mRNA, which includes exons and introns. During the process of RNA splicing, introns are removed and exons joined to form a contiguous coding sequence. This "mature" mRNA is ready for translation.

Exons and introns are related to genes. An exon is termed as a nucleic acid sequence which is represented in the RNA molecule. Introns, on the other hand, are termed as nucleotide sequences seen within the genes which are removed through RNA splicing for generating a mature RNA molecule.

In simple words, exons can be termed as DNA bases which are translated into mRNA. Introns are also DNA bases that are found in-between exons.

Introns and exons were discovered independently by American molecular biologists Richard Roberts and Phillip Sharp in 1977.

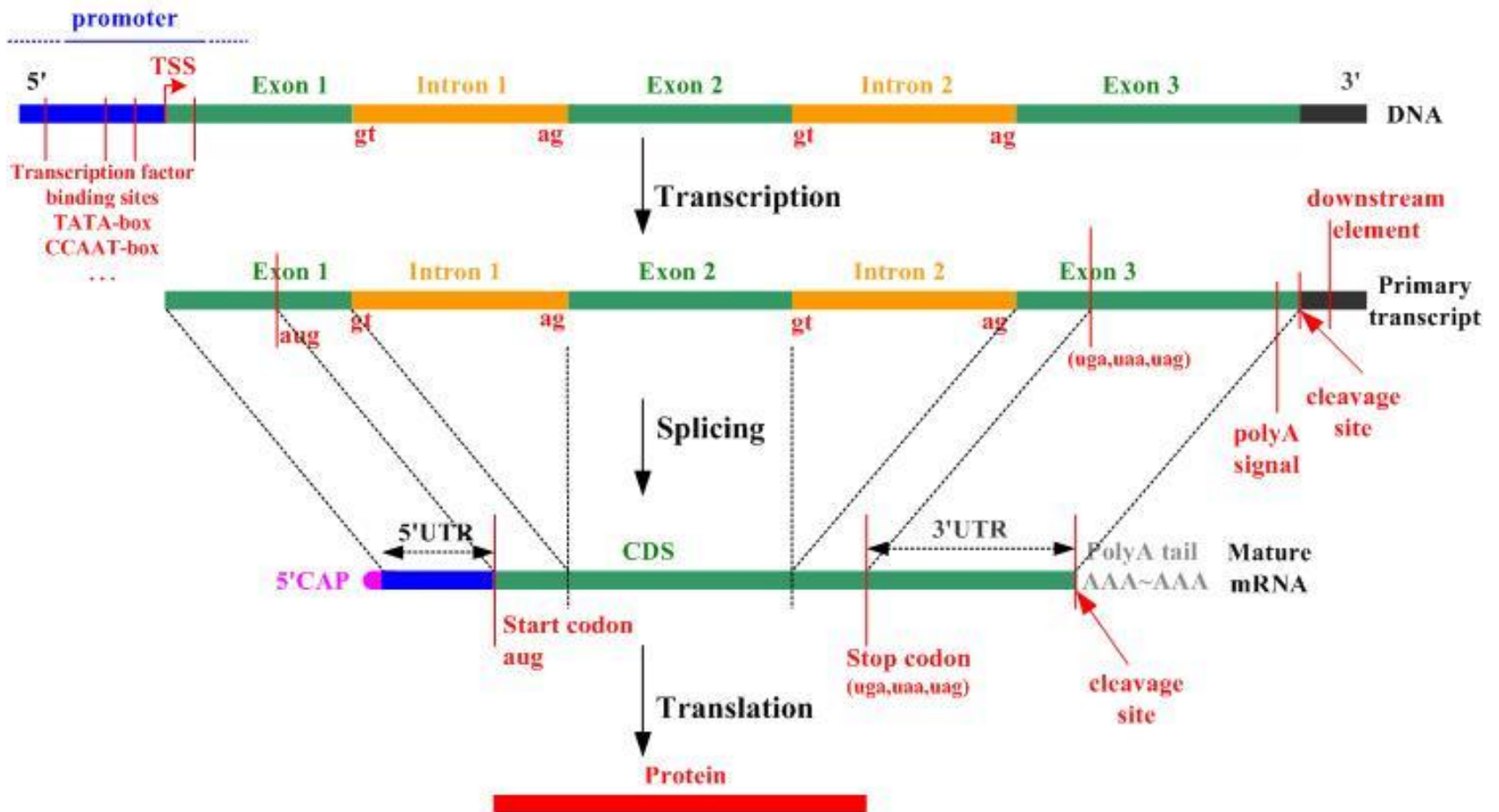
Introns are very much common in the genome of higher vertebrates such as human beings and mice. On the other hand, introns are unlikely to be seen in the genome of certain varieties of eukaryotic micro-organisms such as baker's yeast but are seen in archaeal and bacterial genes.

It can also be seen that introns are less conserved which means that their sequence changes very [frequently](#) over time. On the contrary, exons are very much conserved which means that their sequence does not change rapidly over [time](#) or in-between the species.

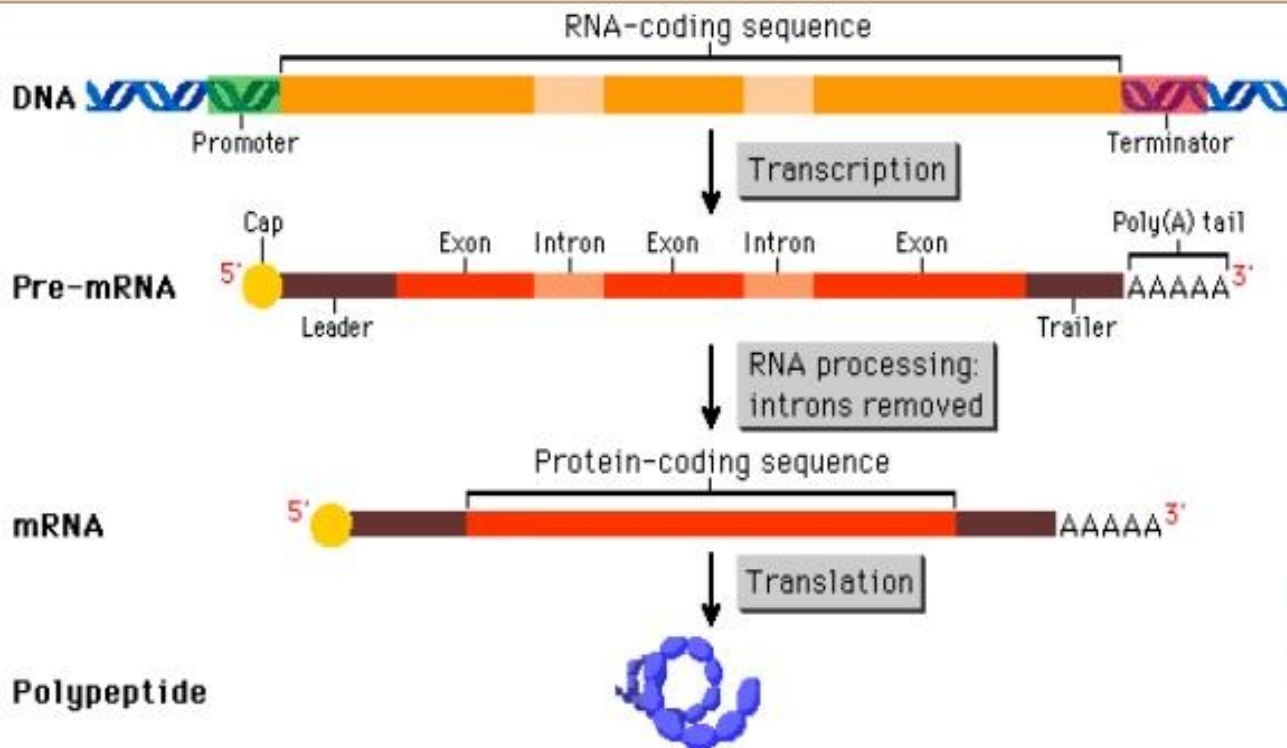
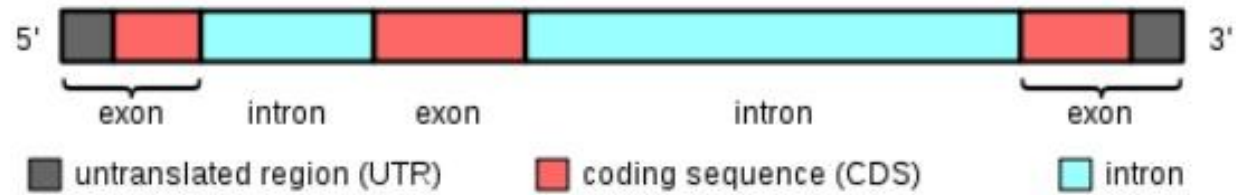
While exons are codes of proteins, introns are not at all implicated with the protein coding. So it can be said that exons are coding areas whereas introns are non-coding areas.

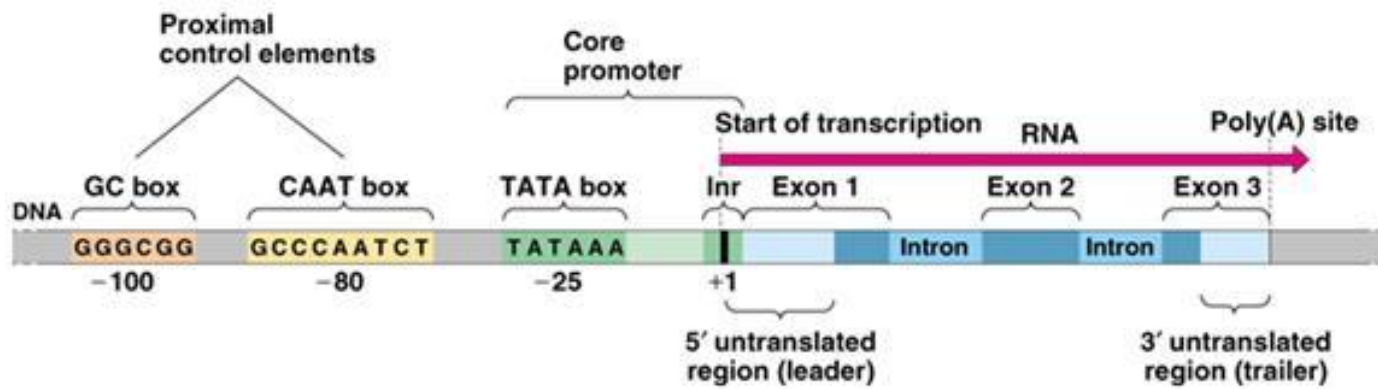
The term 'intron' was derived from 'intrinsic region,' a region inside a gene. Introns are also sometimes termed as intervening sequences. 'Exon' is a term that is derived from 'expressed region.' Walter Gilbert, an American biochemist, coined the term.



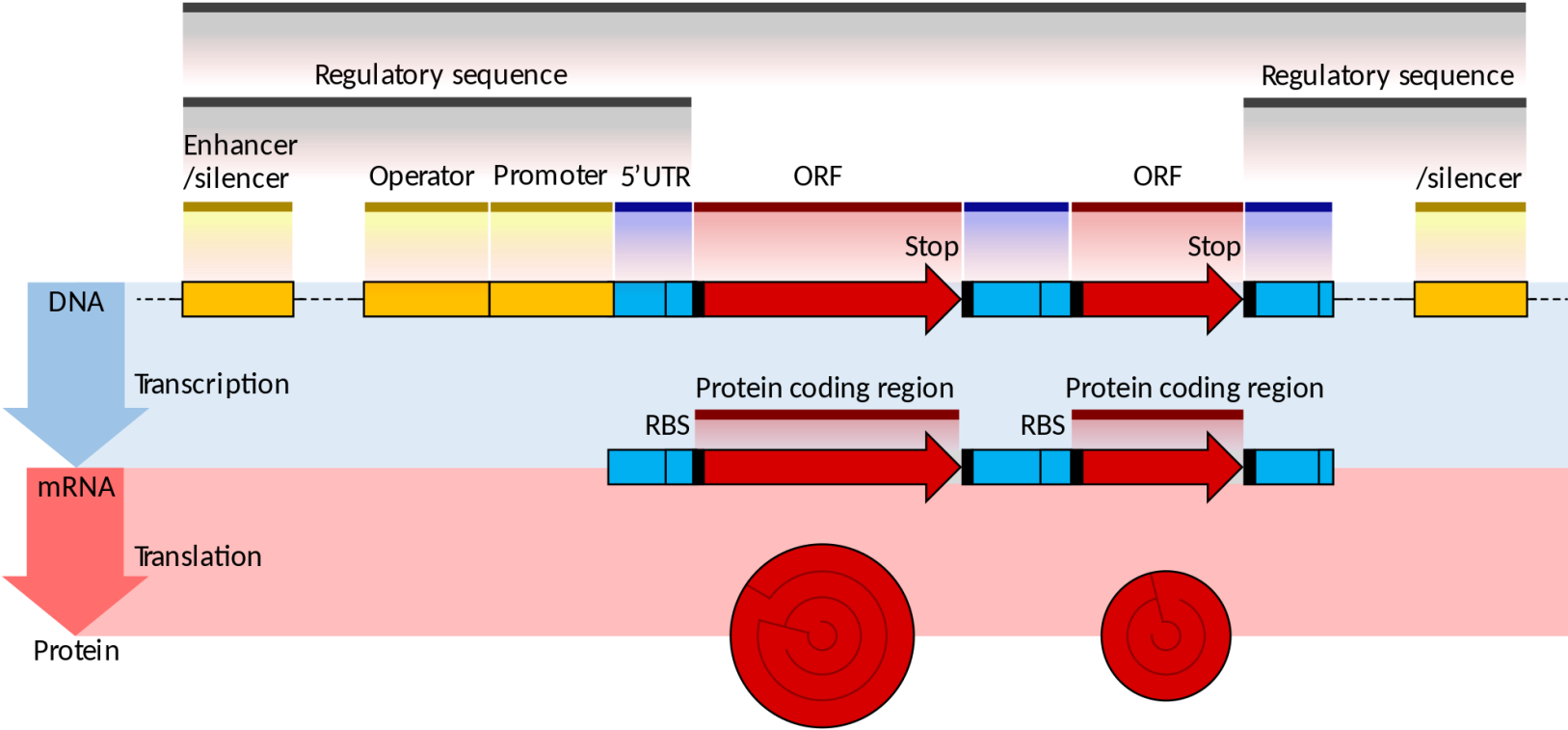


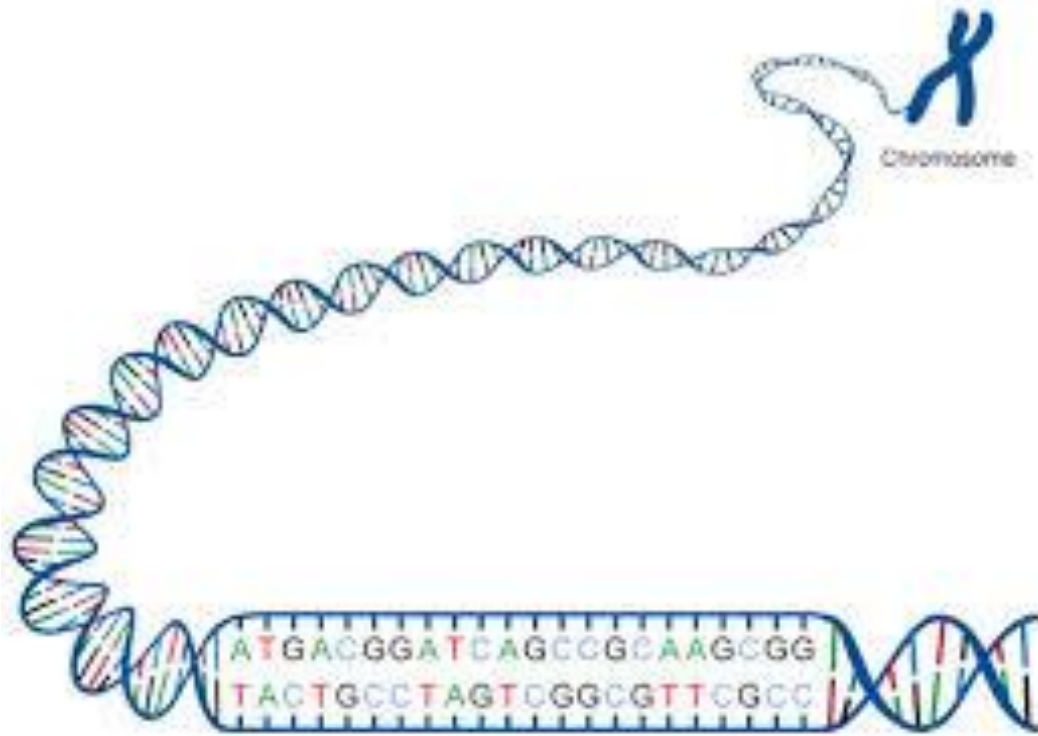
Eukaryotic gene.





Polycistronic operon



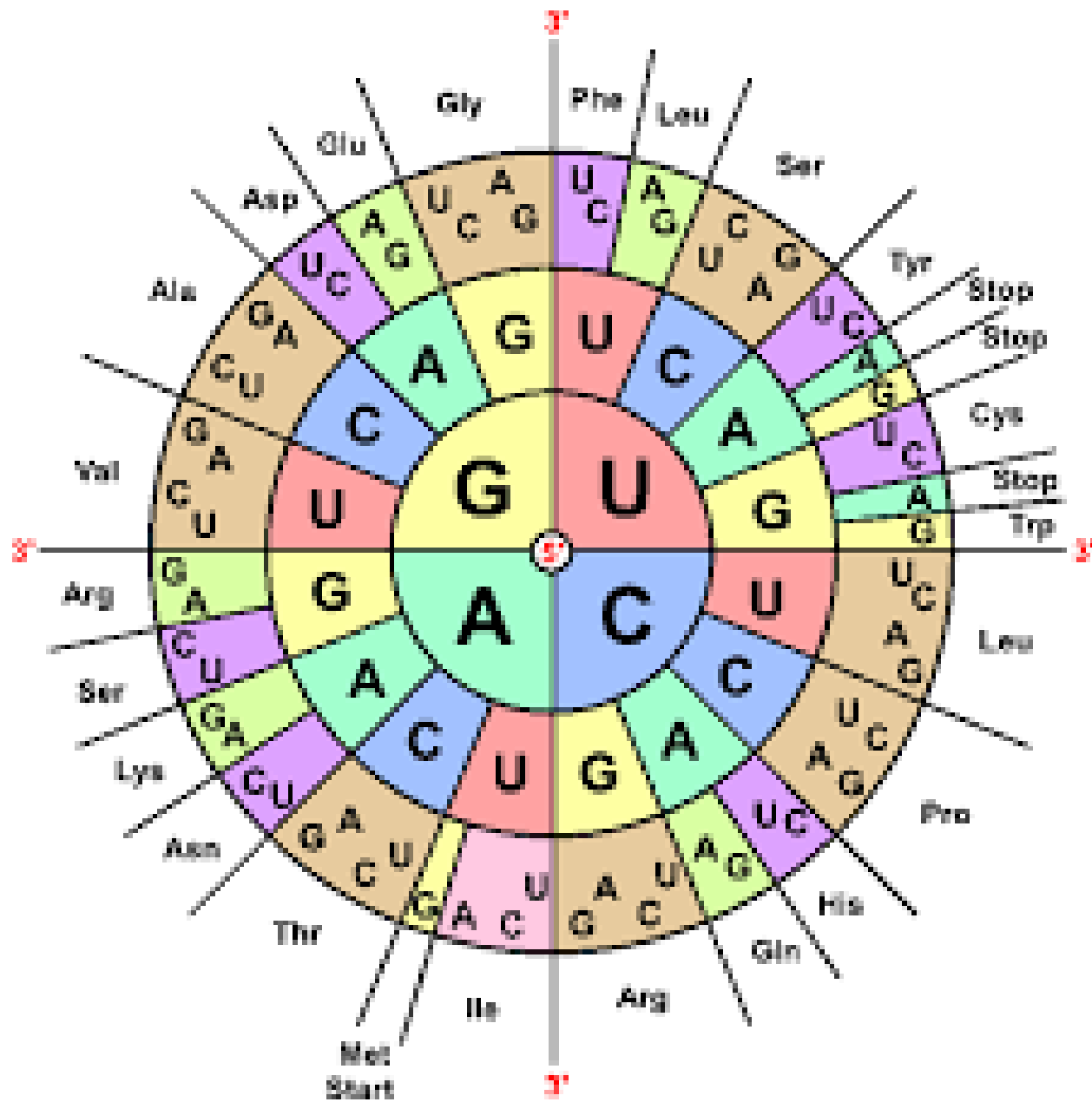


Genetic code is a set of rules by which information encoded in genetic material (DNA or RNA) is translated into protein by living cells.

## What is genetic code?

- The **genetic code** is the set of rules by which information encoded in **genetic material** (DNA or RNA sequences) is translated into proteins by living cells.

5'							
	G	C	A	G			
UUU	Phenylalanine	UUU		UUU	Tyrosine	UUU	Leucine
UUC	Phenylalanine	UUC	Proline	UUC		UUC	Leucine
UUA	Leucine	UUA		UUA	Valine	UUA	Leucine
UUG	Leucine	UUG		UUG	Valine	UUG	Leucine
UUU		UUU		UUU	Valine	UUU	Leucine
UUC		UUC	Proline	UUC		UUC	Leucine
UUA		UUA		UUA	Valine	UUA	Leucine
UUG		UUG		UUG	Valine	UUG	Leucine
UUU		UUU		UUU	Valine	UUU	Leucine
UUC		UUC	Proline	UUC		UUC	Leucine
UUA		UUA		UUA	Valine	UUA	Leucine
UUG		UUG		UUG	Valine	UUG	Leucine
UUU		UUU		UUU	Valine	UUU	Leucine
UUC		UUC	Proline	UUC		UUC	Leucine
UUA		UUA		UUA	Valine	UUA	Leucine
UUG		UUG		UUG	Valine	UUG	Leucine
UUU		UUU		UUU	Valine	UUU	Leucine
UUC		UUC	Proline	UUC		UUC	Leucine
UUA		UUA		UUA	Valine	UUA	Leucine
UUG		UUG		UUG	Valine	UUG	Leucine



4 bases and 3 positions:

$4 \times 4 \times 4 = 64$  Codons

Twenty amino acids (Val, Lys, Arg etc) bind either one/more than one codons

AUG-Met-Start Codon

UAA, UGA and UAG are stop codon

Characteristics:

1. In general, codons are universal in all organism with few exception
2. Use to prepare energy i.e. protein



# Genetic Code- Table

		Second Letter					
		U	C	A	G		
1st letter	U	UUU   Phe UUC   UUA   Leu UUG	UCU   UCC   Ser UCA   UCG	UAU   Tyr UAC   UAA   Stop UAG   Stop	UGU   Cys UGC   UGA   Stop UGG   Trp	U C A G	
	C	CUU   CUC   Leu CUA   CUG	CCU   CCC   Pro CCA   CCG	CAU   His CAC   CAA   Gln CAG	CGU   CGC   Arg CGA   CGG	U C A G	
	A	AUU   AUC   Ile AUA   AUG   Met	ACU   ACC   Thr ACA   ACG	AAU   Asn AAC   AAA   Lys AAG	AGU   Ser AGC   AGA   Arg AGG	U C A G	
	G	GUU   GUC   Val GUA   GUG	GCU   GCC   Ala GCA   GCG	GAU   Asp GAC   GAA   Glu GAG	GGU   GGC   Gly GGA   GGG	U C A G	

3rd

letter

	Isoleucine	Threonine	Lysine	Arginine	A
G	Start Methionine	Threonine	Lysine	Arginine	G
	Valine	Alanine	Aspartic acid	Glycine	U
	Valine	Alanine	Aspartic acid	Glycine	C
	Valine	Alanine	Aspartic acid	Glycine	A
	Valine	Alanine	Aspartic acid	Glycine	G

AUG

GUU

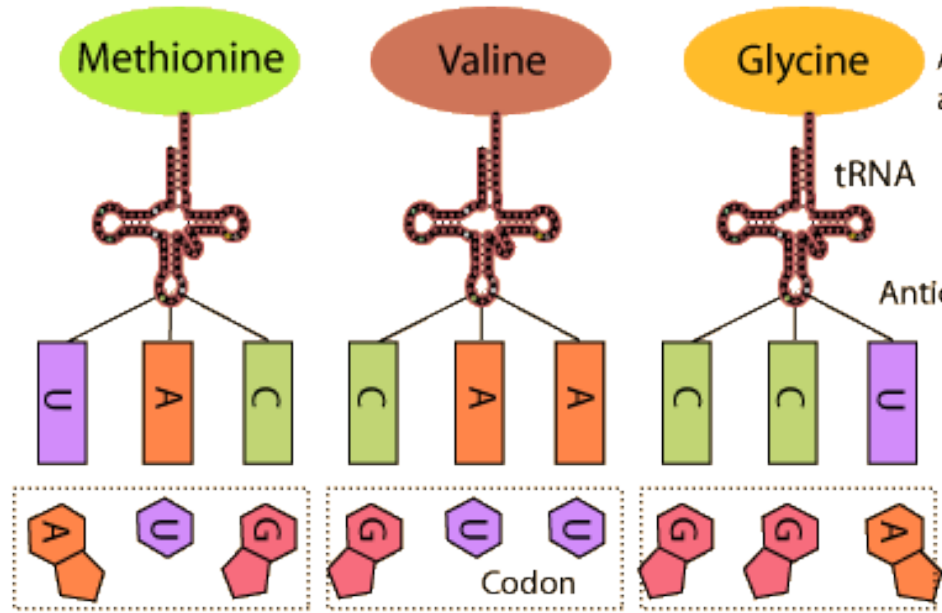
GGA

Methionine

Valine

Glycine

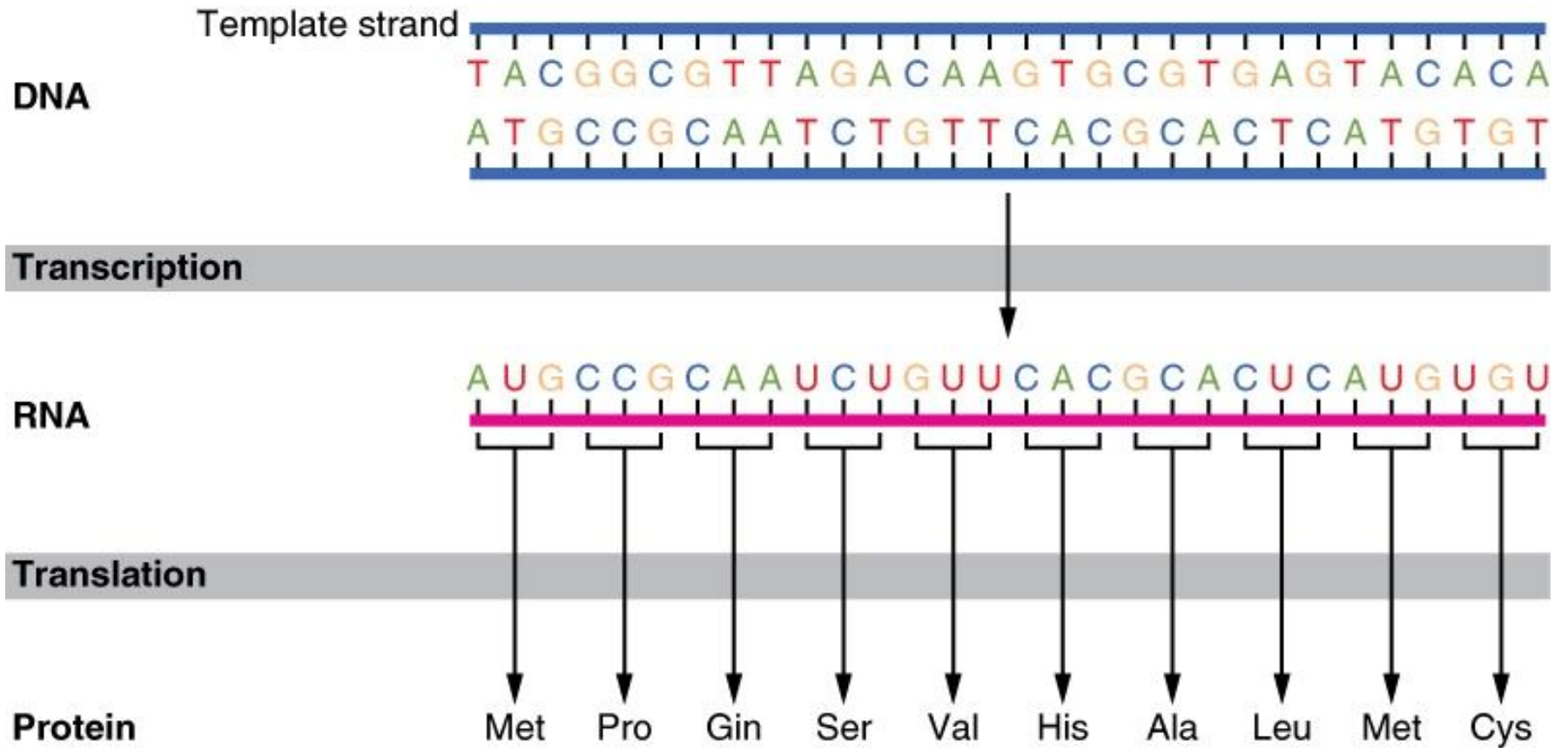
Amino acid



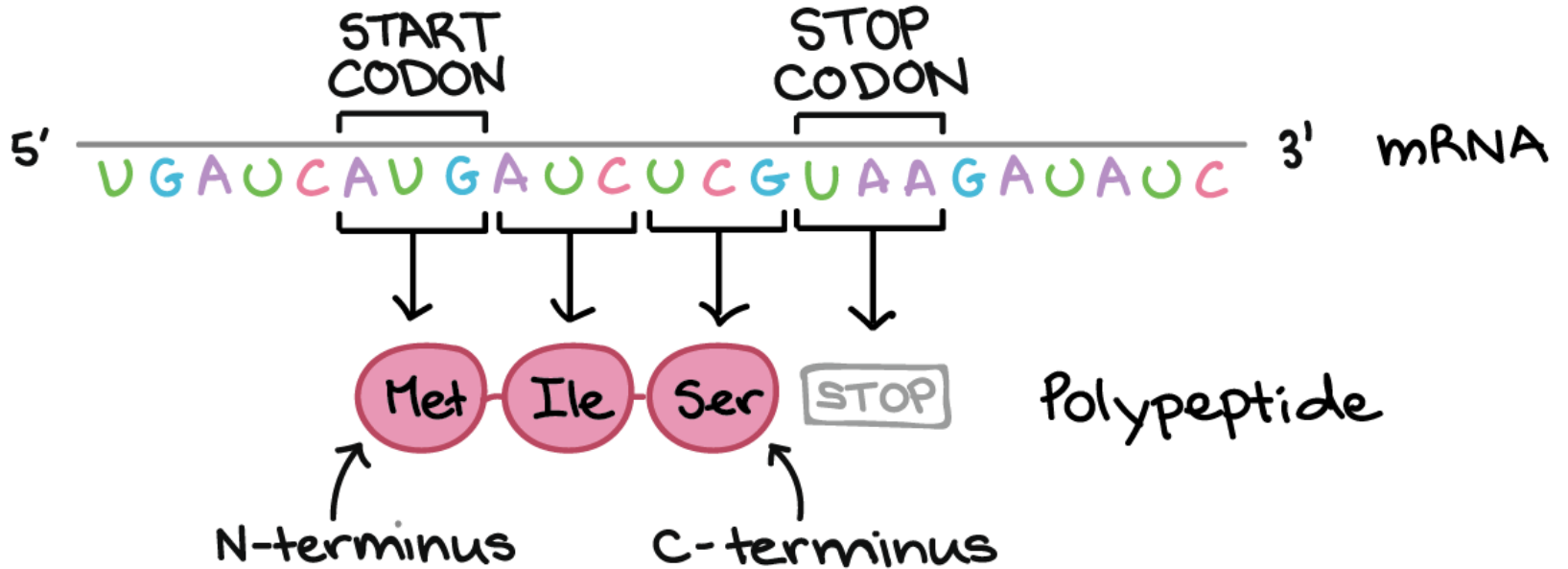
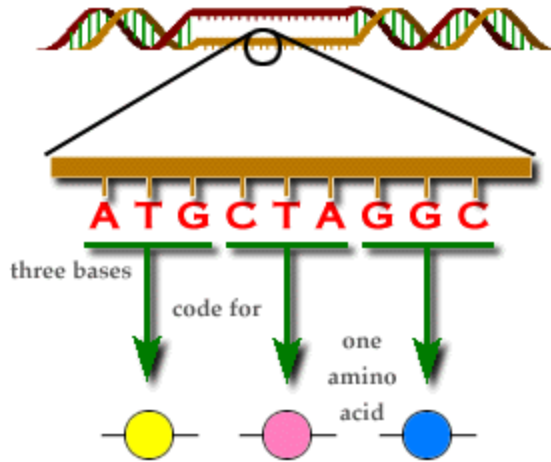
mRNA transcribed code

The amino acid associated with a given triplet of bases does not appear to be chemically determined.

The presence of the anticodon on one end of the tRNA does not chemically determine the amino acid attached to the other end.



# The Genetic Code



# Genetic Code

- The genetic code is the set of rules by which information encoded in genetic material is translated into proteins by living cells.
- The information in DNA is in the form of triplet codons.
- It is first transcribed into RNA and then into proteins.
- Every triplet codon in the DNA specifies one amino acid in the protein.

