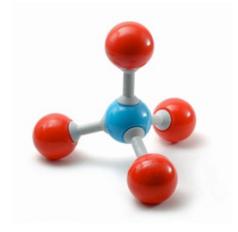
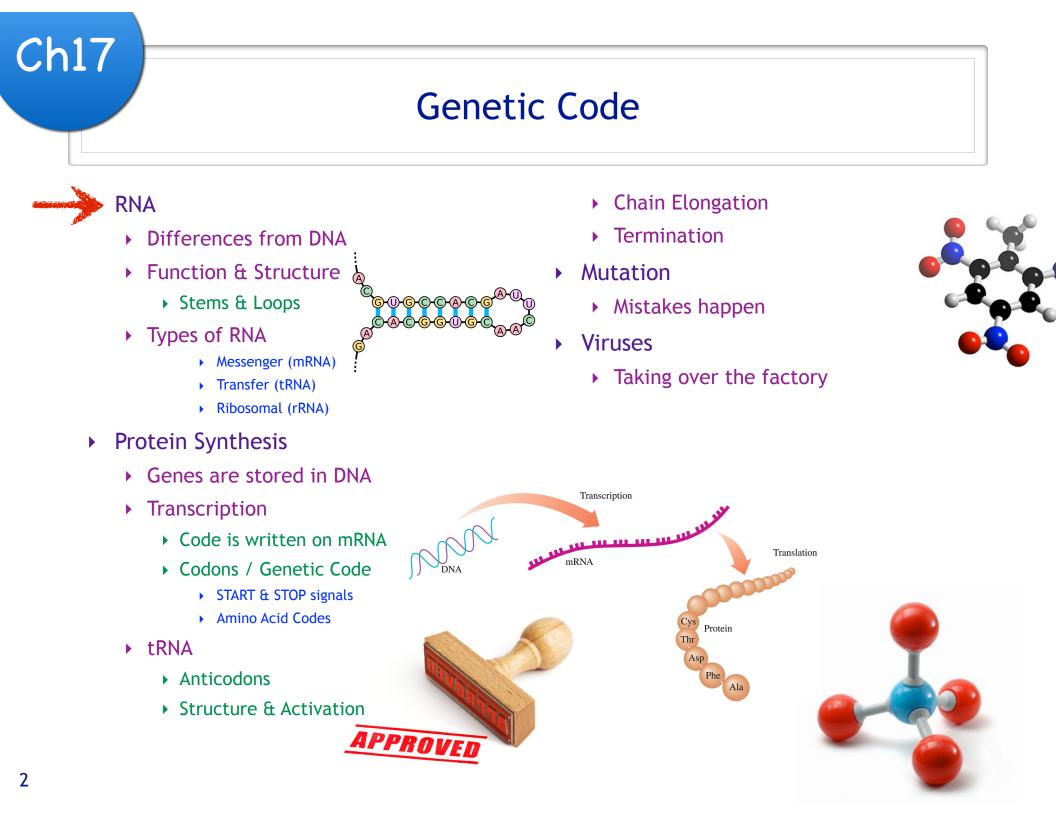


Reading and using the data stored in DNA.



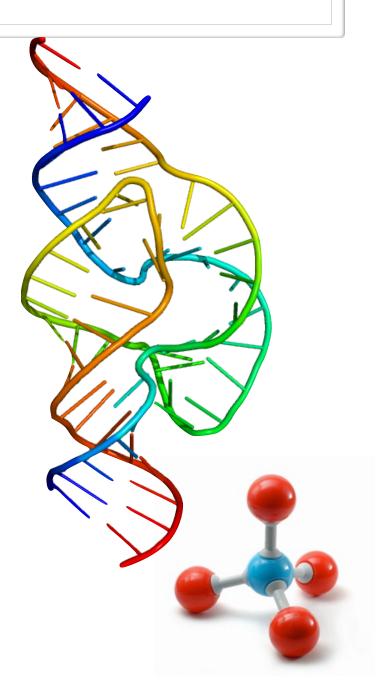
version 1.0 © Nick DeMello, PhD. 2007-2015





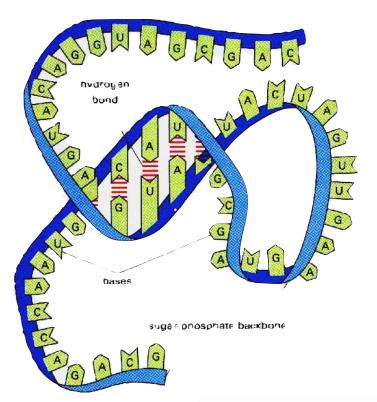
RNA

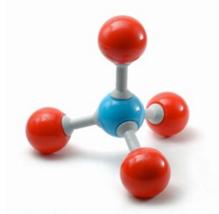
- RNA makes up most of the nucleic acid found in the cell.
- It moves the genetic information needed for cell operation from where it's stored to where it's used.
- RNA molecules are polymers of nucleotides and differ from DNA molecules in four important ways:
 - 1. The sugar in RNA is ribose rather than the deoxyribose found in DNA.
 - 2. The base uracil replaces thymine.
 - 3. RNA molecules are single stranded; DNA is double stranded.
 - 4. RNA molecules are much smaller than DNA molecules.



RNA Structure

- RNA is a smaller molecule than DNA.
- But it's still a big molecule, which can fold over on itself.
- Because it is composed of a single strand it's bases are unpaired.
- The bases can pair with bases elsewhere in the same chain.
- These strong intramolecular interactions between base pairs produce tertiary structures.
- Common tertiary structures include:
 - Stems (intramolecular double helix)
 - Loops



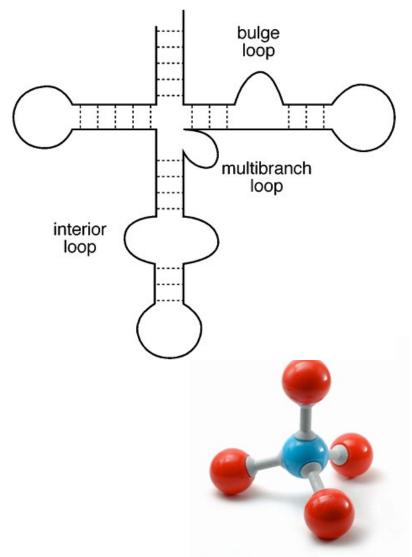


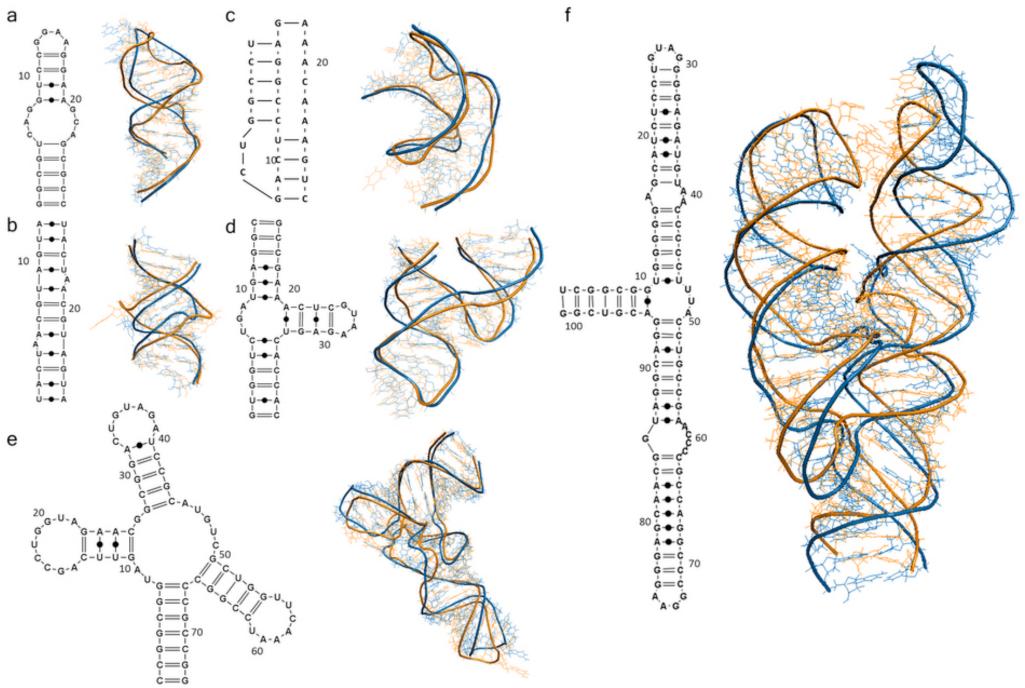
RNA Structure

hairpin

loop

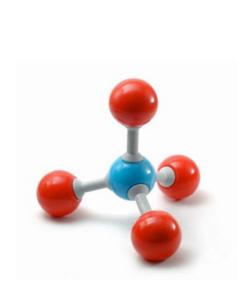
- Loops occur in RNA when the base pairs on either side of the loop form intramolecular bonds
- There are different kinds of loops depending on their location:
 - Bulge loops
 - Interior loops
 - Multibranch loops
 - Stem loops (or hairpin loops)
- The resulting structure is a key building block of many larger RNA secondary structures.





RNA Structure

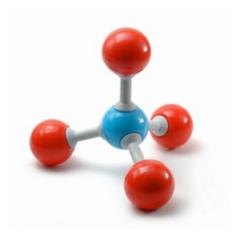
- Loops can occur in single strand DNA but are much more common in RNA.
- Loops that are less than three bases long have too much ring strain and do not form.
- Large loops with no stem are also unstable.
- Optimal loop length is 4-8 bases long.
- One common loop with the sequence UNCG (N= U, A, C, G) is known as a "tetraloop" and is particularly stable due to the base-stacking interactions of its component nucleotides.
- The three most common loops contain the sequences GNRA, UNCG or CUYG.



Three types of RNA

- RNA transmits information from DNA to make proteins and has several types:
- Messenger RNA (mRNA) carries genetic information from DNA in the nucleus to the ribosomes in the cytoplasm.
- Transfer RNA (tRNA) interprets the genetic information in mRNA and brings specific amino acids to the ribosome for protein synthesis.
- Ribosomal RNA (rRNA), the most abundant type of RNA, is combined with proteins to form ribosomes.

| Туре | Abbreviation | Function in the Cell | Percentage of Total RNA |
|---------------|--------------|--|----------------------------|
| Ribosomal RNA | rRNA | Synthesizes protein; major component of the ribosomes | 80 |
| Messenger RNA | mRNA | Carries information for protein synthesis from the DNA in the nucleus to the ribosomes | 5 |
| Transfer RNA | tRNA | Brings amino acids to the ribosomes for protein synthesis | 15 |



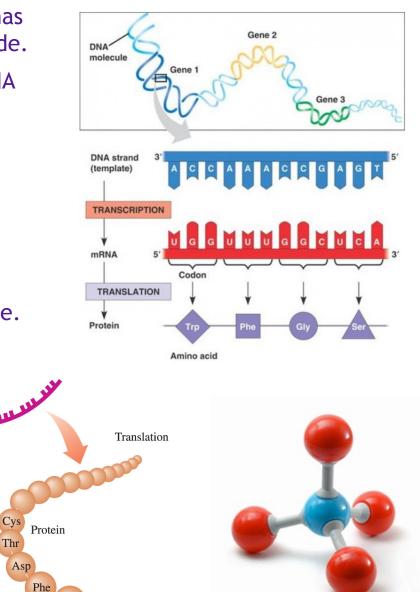
Ch17 **Genetic Code** Chain Elongation ► RNA Termination Differences from DNA Function & Structure **Mutation** Stems & Loops Mistakes happen Types of RNA Viruses Þ Messenger (mRNA) Taking over the factory Transfer (tRNA) Ribosomal (rRNA) **Protein Synthesis** Genes are stored in DNA Transcription Transcription Code is written on mRNA Translation mRNA Codons / Genetic Code ► START & STOP signals Amino Acid Codes Protein Thr ▶ tRNA Anticodons Structure & Activation PPROV 9

Transcription

Ala

mRNA

- A gene is a section of one strand of DNA that has the sequence needed to build a specific peptide.
- That sequence get's copied into messenger RNA (mRNA) by the process of transcription.
- The messenger RNA brings that information to the ribosomes.
- The ribosomes build the amino acid from the sequence by the process of translation.
- We'll look at those two processes one at a time.



 $C \rightarrow G$

 $G \rightarrow C$

 $A \rightarrow U$

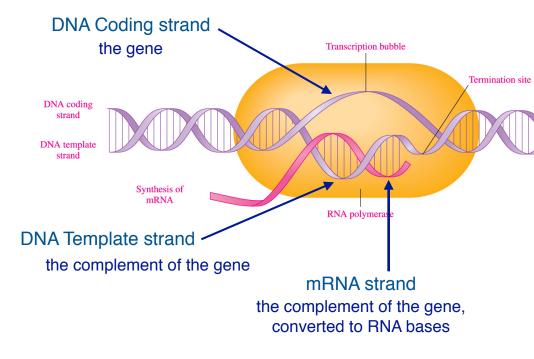
 $T \rightarrow A$

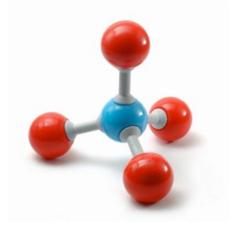
RNA

side

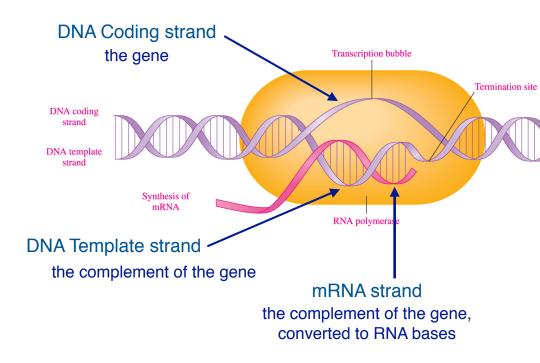
- Transcription involves building a new mRNA molecule based on the gene information in DNA.
- The mRNA is built by the enzyme mRNA polymerize.
- mRNA is synthesized using complementary base pairing
 - Cytosine (C) and guanine (G) pair to each other in DNA and RNA.
 - DNA adenine (A) pairs with RNA uridine (U)
 - DNA thymine (T) paris with RNA adenine (A)

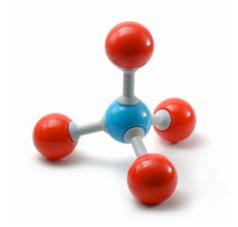
DNA side



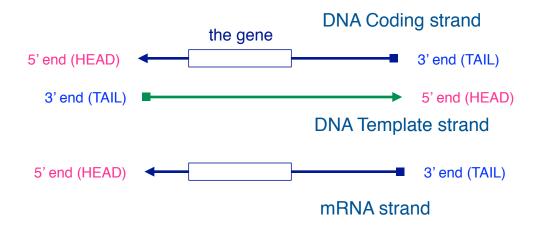


- Transcription involves three strands of nucleic acid.
 - The DNA coding strand.
 - This is the strand which contains the sequence defined as the gene.
 - The DNA template strand.
 - This is the complementary sequence to the gene.
 - mRNA reads from this strand, the one opposite the gene.
 - The mRNA strand.
 - The mRNA strand is built with bases that complement the template strand.



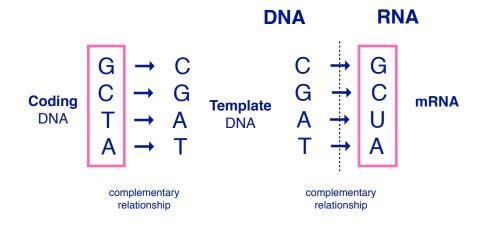


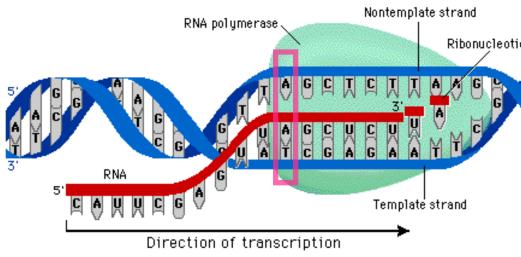
- Each pair of interacting strands run in opposite directions.
 - The DNA coding strand and DNA template strand run head to tail.
 - The 5' end of one matches to the 3' of the other.
 - The DNA template strand and mRNA strand run head to tail.
 - The 5' end of one matches to the 3' of the other.
- That means going from the gene to the mRNA strand, the directions are reversed twice.
- The sequence in the gene and in the mRNA strand run in the same direction.

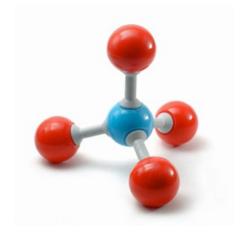




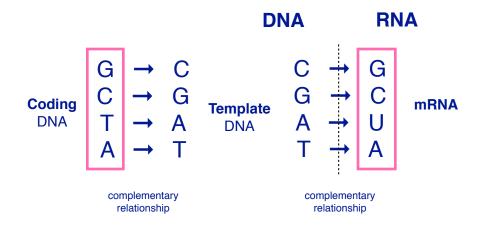
- Because base pairs are complimentary in both processes, the gene expressed in the mRNA is the same as the gene expressed in the coding DNA.
 - EXCEPT: <u>T in DNA becomes U in RNA</u>







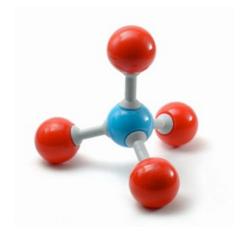
- Because base pairs are complimentary in both processes, the gene expressed in the mRNA is the same as the gene expressed in the coding DNA.
 - EXCEPT: T in DNA becomes U in RNA



DNA Coding strand

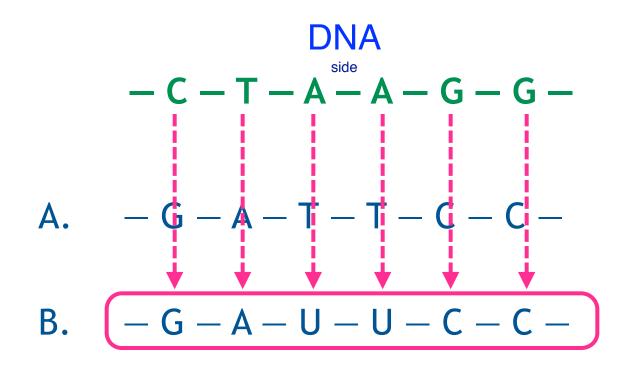
3'
$$- T - C - G - A - G - 5'$$

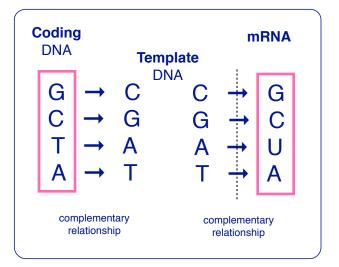
mRNA strand 5'
$$-A - G - C - U - C - 3'$$



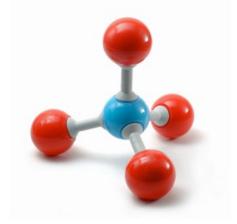
Try it.

• Given this section of template DNA what sequence of mRNA is produced?



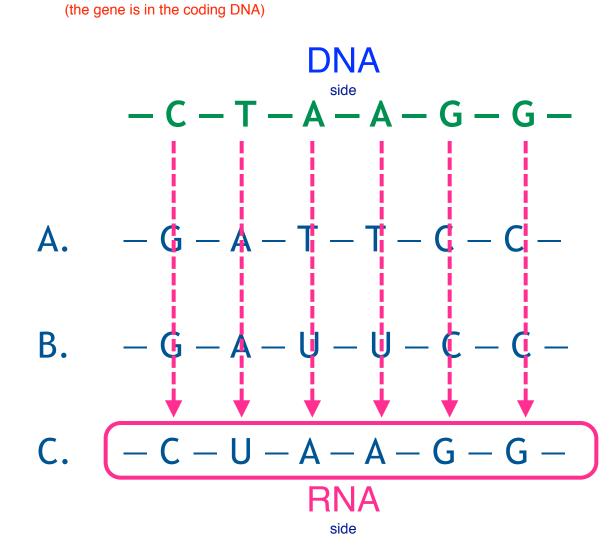


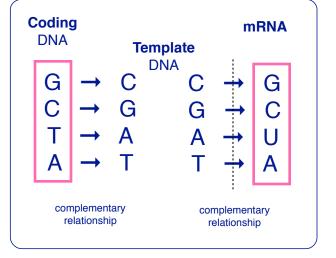
C.
$$-C - U - A - A - G - G - RNA_{side}$$

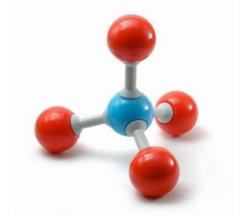


Try it.

• Given this Gene what sequence of mRNA is produced?







Ch17 **Genetic Code** Chain Elongation ► RNA Termination Differences from DNA Function & Structure **Mutation** Stems & Loops Mistakes happen Types of RNA Viruses Messenger (mRNA) Taking over the factory Transfer (tRNA) Ribosomal (rRNA) Protein Synthesis Genes are stored in DNA Transcription Transcription Code is written on mRNA Translation mRNA Codons / Genetic Code ▶ START & STOP signals Amino Acid Codes Protein Thr ▶ tRNA Anticodons Structure & Activation PPROL

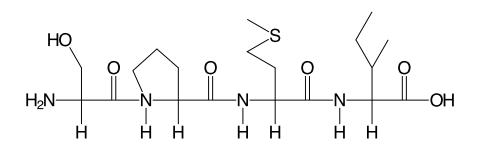
Reading Genetic Code

- Three nucleotides in a row is a triplet.
- Some combinations of nucleotides in a triplet specify an amino acid or instruction.
 - Codons are combinations that mean specific actions or amino acids.
- Codons of genetic code are like bytes of computer code.

a computer says "01001000" → this means "H" a computer says "01100101" → this means "e" "01001000011001010110110001101101101111" is "Hello"

> mRNA says "UCA" \rightarrow this means Serine mRNA says "CCG" \rightarrow this means Proline

"UCACCGAUGAUA" is "Serine Proline Methionine Isoleucine "



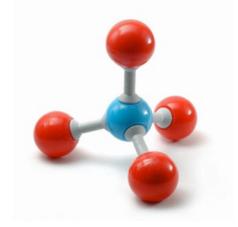
| Binary Bits | Character |
|--------------------|-----------|
| 01001000 | Н |
| 01100101 | e |
| 01101100 | 1 |
| 01101100 | 1 |
| 01101111 | 0 |
| 00100000 | space _ |
| 01001010 | J |
| 01101111 | 0 |
| 01110011 | S |
| 01101000 | h |
| | |

An amino acid can have more than one codon: UCA means Serine UCC also means Serine

Reading Genetic Code

- Three nucleotides in a row is a triplet.
- Some combinations of nucleotides in a triplet specify an amino acid or instruction.
 - Codons are combinations that mean specific actions or amino acids.
- Other codons are instructions on how to use those amino acids.
 - The beginning of a mRNA sequence is called the 5' UTR (untranslated region).
 - The UTR has some information specific to different kinds of cell processes.
 - At some point a AUG byte is introduced. AUG means Methionine.
 - The first methionine encountered tells the ribosomes to start building with the next codon.
 - This is the START codon.
 - What follows is the blue print, the sequence or primary structure, for a specific protein.
 - When that's complete the file is ended with either UAA, UAG, or UGA.
 - This is the STOP codon.

| Binary Bits | Character |
|-------------|-----------|
| 01001000 | Н |
| 01100101 | e |
| 01101100 | 1 |
| 01101100 | 1 |
| 01101111 | 0 |
| 00100000 | space - |
| 01001010 | J |
| 01101111 | 0 |
| 01110011 | S |
| 01101000 | h |
| 01110101 | u |
| 01100001 | a |
| | |



Reading Genetic Code

- A total of 64 codons are possible from the triplet combinations of A, G, C, and U.
- Codons have been determined for all 20 amino acids needed to build a protein.
- At the beginning of mRNA, the codon AUG signals the start of protein synthesis.
- A codon stop signal of UGA, UAA, and UAG in mRNA signals for the termination of protein synthesis.

| mRNA Codon to Amino Acid Translation |
|--------------------------------------|
|--------------------------------------|

| | U | | С | | Α | | G | |
|---|-----|---------|-----------------|---------|-----|--------------------|-----|----------|
| | UUU | | UCU | | UAU | | UGU | 0)(0,(0) |
| U | UUC | PHE (F) | UCC | | UAC | TYR (Y) | UGC | CYS (C) |
| U | UUA | | UCA | SER (S) | UAA | STOP | UGA | STOP |
| | UUG | LEU (L) | UCG | | UAG | 510P | UGG | TRP (W) |
| | CUU | | CCU | CU | CAU | | CGU | |
| 6 | CUC | | CCC | | CAC | HIS (H) GLN (Q) | CGC | ARG (R) |
| C | CUA | LEU (L) | CCA | PRO (P) | CAA | | CGA | |
| | CUG | | CCG | - | CAG | | CGG | |
| | AUU | | ACU | | AAU | | AGU | |
| | AUC | ILE (I) | ACC | | AAC | ASN (N) | AGC | SER (S) |
| Α | AUA | | ACA THR (T) AAA | | AGA | | | |
| | AUG | MET (M) | ACG | | AAG | LYS (K) | AGG | ARG (R) |
| | GUU | | GCU | | GAU | | GGU | |
| 6 | GUC | | GCC | ALA (A) | GAC | ASP (D) | GGC | |
| G | GUA | VAL (V) | GCA | | GAA | | GGA | GLY (G) |
| | GUG | | GCG | | GAG | GLU (E) | GGG | |

This table will be provided on exams and quizzes.

Try it.

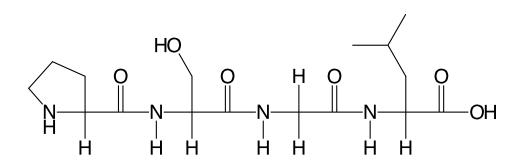
• Determine the amino acids from the following codons in a section of mRNA.

- CCUAGCGGACUU -

- CCU - AGC - GGA - CUU -

Pro Ser Gly Leu

$$-$$
 Pro $-$ Ser $-$ Gly $-$ Leu $-$



mRNA Codon to Amino Acid Translation

| | U | | С | | Α | | G | |
|---|-----|---------|---------|---------|-----|---------|-----|---------|
| | UUU | PHE (F) | UCU | | UAU | | UGU | |
| | UUC | | UCC | | UAC | TYR (Y) | UGC | CYS (C) |
| U | UUA | LEU (L) | UCA | SER (S) | UAA | STOP | UGA | STOP |
| | UUG | LEU (L) | UCG | | UAG | 3106 | UGG | TRP (W) |
| | CUU | | CCU | | CAU | HIS (H) | CGU | |
| c | CUC | | CCC PRO | | CAC | піз (п) | CGC | ARG (R) |
| | CUA | LEU (L) | | | CAA | GLN (Q) | CGA | |
| | CUG | | CCG | | CAG | | CGG | |
| | AUU | | ACU | | AAU | ASN (N) | AGU | SER (S) |
| | AUC | ILE (I) | ACC | | AAC | | AGC | |
| A | AUA | | ACA | THR (T) | AAA | | AGA | |
| | AUG | MET (M) | ACG | | AAG | LYS (K) | AGG | ARG (R) |
| | GUU | | GCU | | GAU | | GGU | |
| | GUC | | GCC | ALA (A) | GAC | ASP (D) | GGC | |
| G | GUA | VAL (V) | GCA | | GAA | | GGA | GLY (G) |
| | GUG | | GCG | | GAG | GLU (E) | GGG | |

CCU = proline

AGC = serine

GGA = glycine



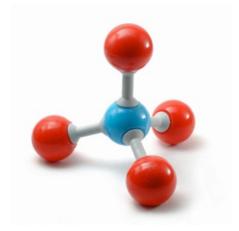
Try it.

• Given the following DNA sequence, what is the first amino acid in the gene?

GGGACAGAATGTGTTGCAGGAACTTC TTCTGGAAGACCTTCTCCTCCTGCAA ATAAAACCTCACCCATGAATGCTCACG CAAGTTTAATTACAGACCTGAA

| | U | | С | | Α | | G | |
|---|-----|---------|-----|---------|---------|---------|-----|---------|
| | UUU | PHE (F) | UCU | | UAU | TYR (Y) | UGU | |
| U | UUC | | UCC | | UAC | 110(1) | UGC | CYS (C) |
| | UUA | LEU (L) | UCA | SER (S) | UAA | STOP | UGA | STOP |
| | UUG | LEO (L) | UCG | | UAG | 3105 | UGG | TRP (W) |
| | CUU | | CCU | CAU | HIS (H) | CGU | | |
| | CUC | | CCC | | CAC | піз (п) | CGC | ARG (R) |
| C | CUA | LEU (L) | CCA | PRO (P) | CAA | GLN (Q) | CGA | |
| | CUG | | CCG | | CAG | | CGG | |
| | AUU | | ACU | | AAU | ASN (N) | AGU | SER (S) |
| | AUC | ILE (I) | ACC | | AAC | | AGC | |
| A | AUA | | ACA | THR (T) | AAA | | AGA | |
| | AUG | MET (M) | ACG | | AAG | LYS (K) | AGG | ARG (R) |
| | GUU | | GCU | | GAU | | GGU | |
| | GUC | | GCC | | GAC | ASP (D) | GGC | |
| G | GUA | VAL (V) | GCA | ALA (A) | GAA | GLU (E) | GGA | GLY (G) |
| | GUG | | GCG | | GAG | GLU (E) | GGG | |

mRNA Codon to Amino Acid Translation



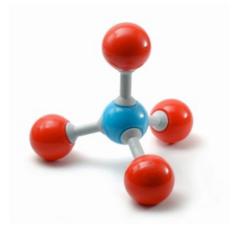
• Given the following DNA sequence, what is the first amino acid in the gene?

GGGACAGAATGTGTTGCAGGAACTTC TTCTGGAAGACCTTCTCCTCCTGCAA ATAAAACCTCACCCATGAATGCTCACG CAAGTTTAATTACAGACCTGAA

start code in RNA is 'AUG' start code in the coding strand of DNA is 'ATG' the first codon in DNA after that is 'TGT' this corresponds to the RNA codon 'UGU' this is the codon for Cysteine.

| | l | J | (| 2 | A | | G | | |
|---|-----|----------|-----|---------|-----|---------|-----|--------------------|--|
| | UUU | PHE (F) | UCU | | UAU | TYR (Y) | UGU | CYS (C) | |
| | UUC | FTIE (F) | UCC | | UAC | 116(1) | UGC | 013(0) | |
| U | UUA | | UCA | SER (S) | UAA | STOP | UGA | STOP | |
| | UUG | LEU (L) | UCG | 1 | UAG | 310F | UGG | TRP (W) | |
| | CUU | | CCU | PRO (P) | CAU | | CGU | | |
| | CUC | | CCC | | CAC | HIS (H) | CGC | ARG (R) | |
| C | CUA | LEU (L) | CCA | | CAA | | CGA | | |
| | CUG | | CCG | | CAG | GLN (Q) | CGG | | |
| | AUU | | ACU | | AAU | ASN (N) | AGU | SER (S) ARG (R) | |
| | AUC | ILE (I) | ACC | | AAC | | AGC | | |
| A | AUA | | ACA | THR (T) | AAA | | AGA | | |
| | AUG | MET (M) | ACG | | AAG | | AGG | | |
| | GUU | | GCU | ALA (A) | GAU | | GGU | | |
| | GUC | VAL (V) | GCC | | GAC | ASP (D) | GGC | | |
| G | GUA | | GCA | | GAA | | GGA | GLY (G) | |
| | GUG | | GCG | | GAG | GLU (E) | GGG | | |

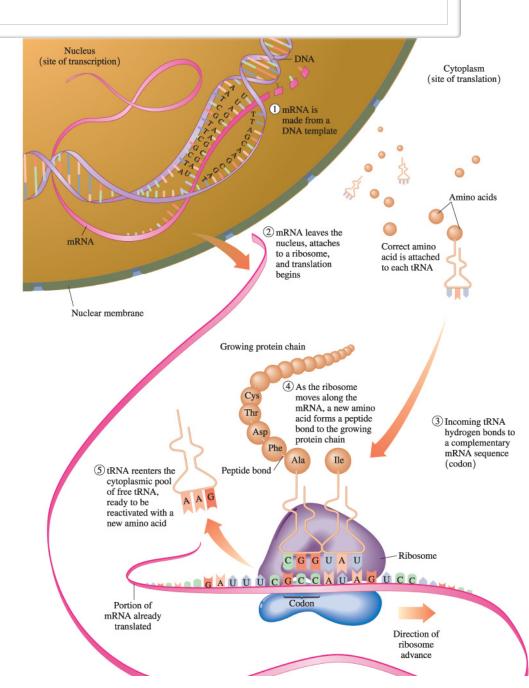
mRNA Codon to Amino Acid Translation



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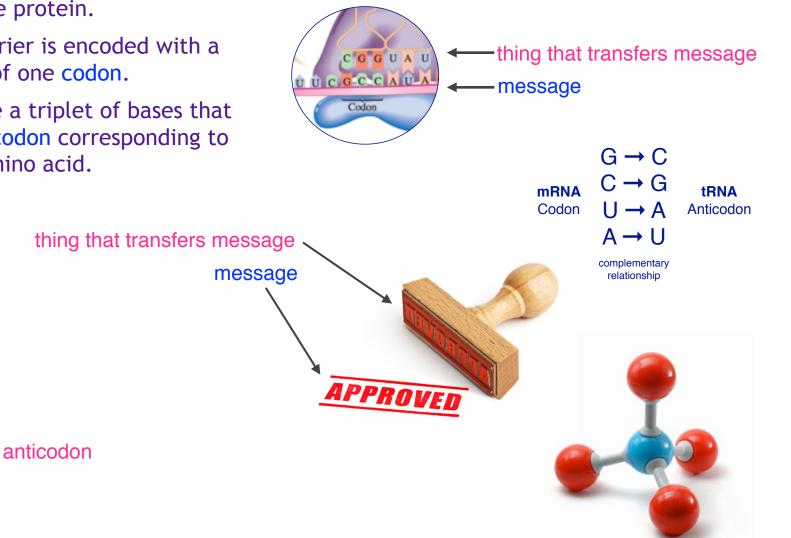
tRNA Translates and Transfers

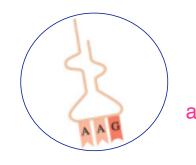
- Translation is the second coding.
- In translation ribosomes make peptide bonds to build the protein.
- tRNA are RNA strands that translate between genetic code amino acids.
- tRNA collects the correct amino acid and delivers it for linking into the protein.



tRNA Translates and Transfers

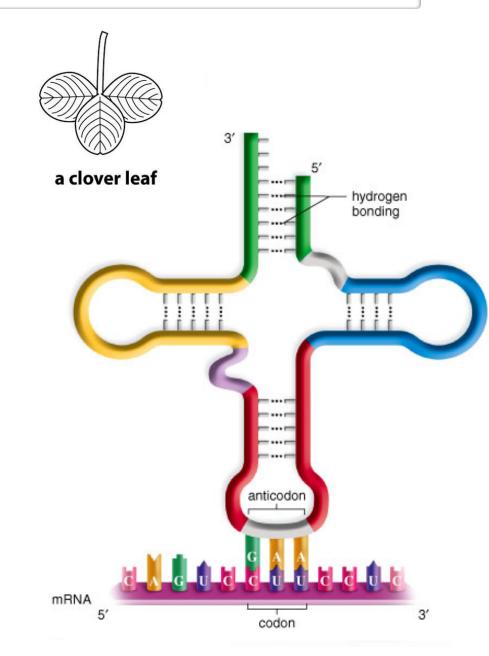
- Ribosomes use tRNA to collect the correct amino acid and deliver it for linking into the protein.
- Each tRNA courier is encoded with a mirror image of one codon.
- Anticodons are a triplet of bases that pair with the codon corresponding to the needed amino acid.





tRNA Translates and Transfers

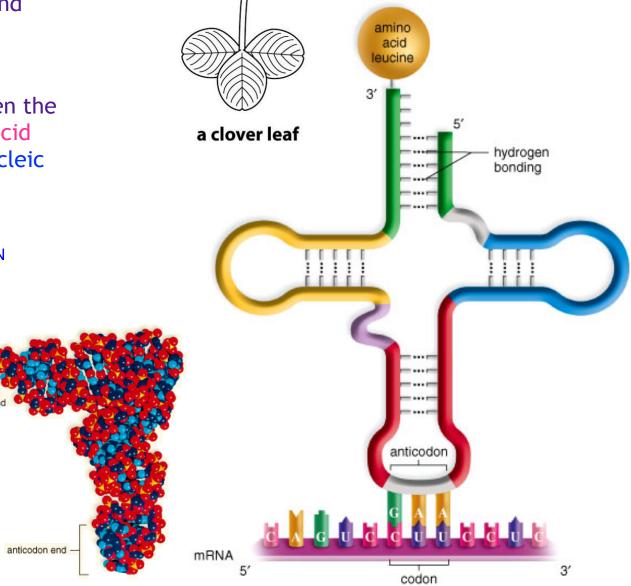
- tRNA Molecules
 - Are of uniform size, 70 to 90 nucleotides.
 - Have a 3'-end that starts with the nucleotide sequence ACC.
 - ACC is known as the acceptor stem.
 - Have an anticodon on a loop opposite the acceptor stem.
 - Have three loops, a clover leaf shape.



Activation of tRNA

tRNA are activated when they bond with an amino acid. tRNA forms an ester bond between the carboxylate group of the amino acid and the hydroxyl group of the nucleic acid on the acceptor stem. $\int_{f}^{h_{p}} \int_{f}^{h_{p}} \int_{f$

amino acid

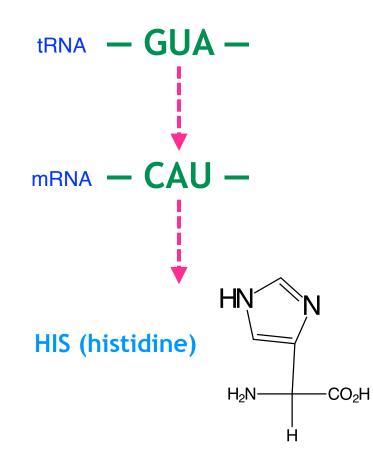


HO

ŃH₂

Try it.

What amino acid does the anticodon GUA identify?



| | U | | С | | Α | | G | |
|---|-----|---------|-----|---------|-----|---------|-----|---------|
| | UUU | | UCU | | UAU | | UGU | |
| | UUC | PHE (F) | UCC | | UAC | TYR (Y) | UGC | CYS (C) |
| U | UUA | | UCA | SER (S) | UAA | STOP | UGA | STOP |
| | UUG | LEU (L) | UCG | | UAG | 310F | UGG | TRP (W) |
| | CUU | | CCU | | CAU | HIS (H) | CGU | |
| | CUC | | CCC | | CAC | піз (п) | CGC | ARG (R) |
| C | CUA | LEU (L) | CCA | PRO (P) | CAA | GLN (Q) | CGA | |
| | CUG | | CCG | | CAG | | CGG | |
| | AUU | | ACU | | AAU | ASN (N) | AGU | SER (S) |
| | AUC | ILE (I) | ACC | | AAC | | AGC | |
| A | AUA | | ACA | THR (T) | AAA | | AGA | |
| | AUG | MET (M) | ACG | | AAG | LYS (K) | AGG | ARG (R) |
| | GUU | | GCU | | GAU | | GGU | |
| G | GUC | | GCC | | GAC | ASP (D) | GGC | GLY (G) |
| G | GUA | VAL (V) | GCA | ALA (A) | GAA | GLU (E) | GGA | |
| | GUG | | GCG | | GAG | | GGG | - |

mRNA Codon to Amino Acid Translation

 $\begin{array}{c} G \rightarrow C \\ \\ \mathsf{mRNA} & C \rightarrow G \\ \\ Codon & U \rightarrow A \\ & A \rightarrow U \end{array}$

complementary relationship



Ch17 Genetic Code **Chain Elongation** ► RNA Termination Differences from DNA Function & Structure **Mutation** Stems & Loops Mistakes happen Types of RNA Viruses Þ Messenger (mRNA) Taking over the factory Transfer (tRNA) Ribosomal (rRNA) Protein Synthesis Genes are stored in DNA Transcription Transcription Code is written on mRNA Translation mRNA Codons / Genetic Code ► START & STOP signals Amino Acid Codes Protein Thr ▶ tRNA Anticodons Structure & Activation PPROV

Protein Synthesis: Translation

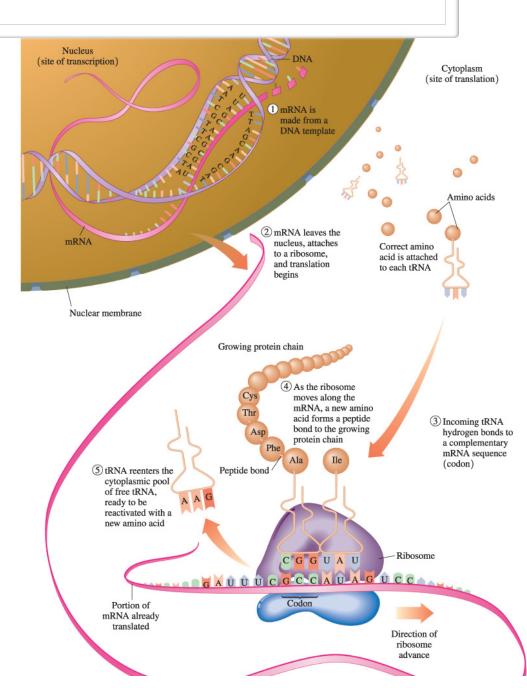
- Translation is the second stage.
- In translation ribosomes make peptide bonds to build the protein.
- tRNA are RNA strands that translate between genetic code amino acids.

Step

1. DNA transcription

- 2. Activation of tRNA
- **3.** Initiation and chain elongation

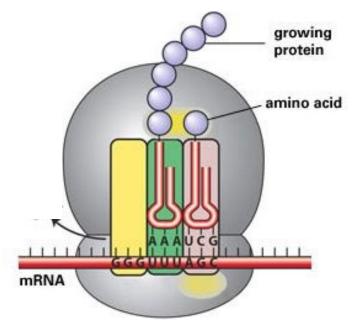
4. Chain termination

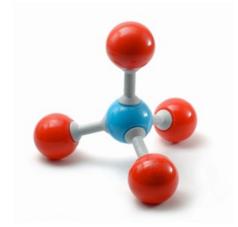


Initiation and Chain Elongation

Protein synthesis begins when mRNA binds to a ribosome.

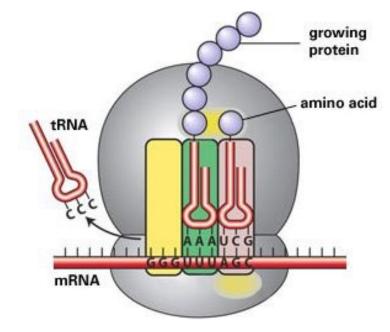
- The first codon in mRNA is a start codon, AUG, which forms hydrogen bonds to methionine-tRNA.
- Another tRNA hydrogen bonds to the next codon, placing a second amino acid adjacent to methionine.
- A peptide bond forms between the C-terminal of methionine and the N-terminal of the second amino acid.

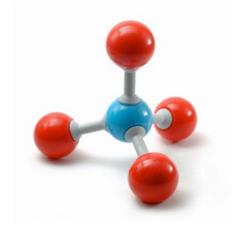




Translocation and Chain Elongation

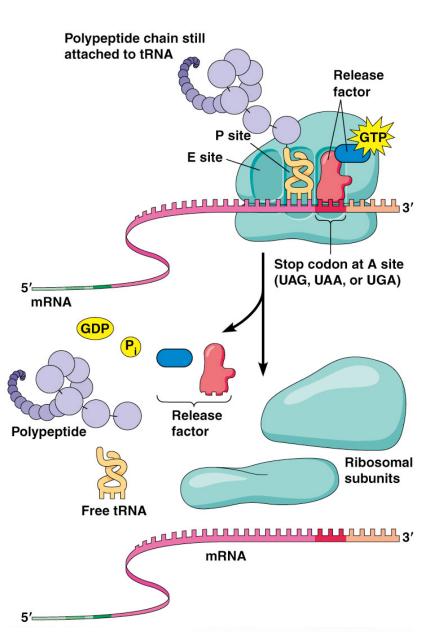
- The initial tRNA detaches from the ribosome, which shifts to the next available codon, a process called translocation.
- During chain elongation, the ribosome moves along the mRNA from codon to codon, so that the tRNAs can attach new amino acids to the growing protein chain.





Chain Termination

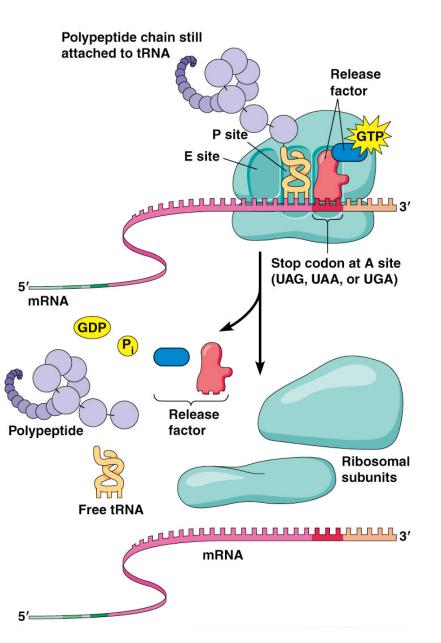
- In the termination step,
 - all the amino acids are linked
 - the ribosome reaches a "stop" codon: UGA, UAA, or UAG
 - there is no tRNA with an anticodon for the "stop" codons
 - the polypeptide detaches from the ribosome
 - the initial amino acid, methionine, is usually removed from the beginning of the protein chain



Termination

Once the polypeptide is released,

- the R groups of the amino acids in the new polypeptide can form hydrogen bonds to give the secondary structures of α helices, β-pleated sheets, or triple helices
- chains form cross-links such as salt bridges and disulfide bonds to produce tertiary and quaternary structures, making it a biologically active protein

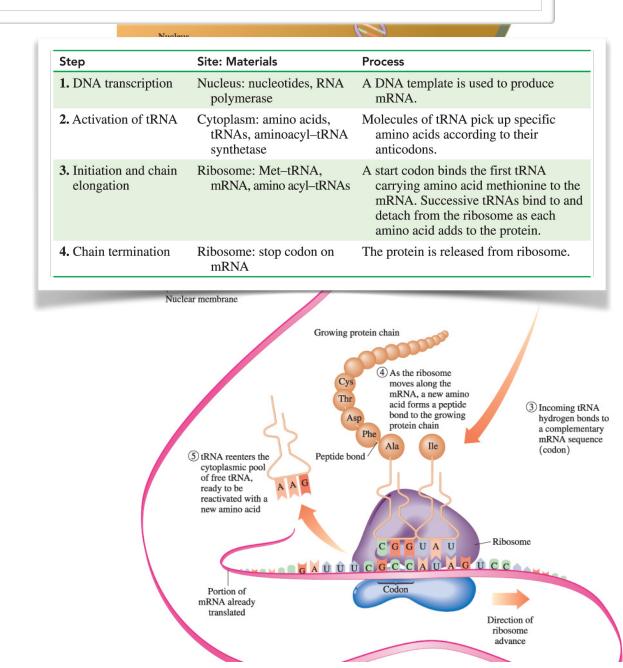


Protein Synthesis Overview

- The DNA strand opens.
 - 1. The template strand gets transcribed to mRNA.
 - 2. tRNA is activated, it picks up the necessary amino acids.

Anticodons on the tRNA translate the mRNA into amino acids.

- 3. Ribosome makes the peptide bonds from amino acids tRNA provides.
- 4. The system finishes and releases the new peptide.



Protein Synthesis Overview

 $C \rightarrow G$

- The DNA strand opens.
 - 1. The template strand gets $G \rightarrow C$ transcribed to mRNA. $T \rightarrow A$
 - 2. tRNA is activated, it picks up the necessary amino acids.

Anticodons on the tRNA translate the mRNA into amino acids.

- 3. Ribosome makes the peptide bonds from amino acids tRNA provides.
- 4. The system finishes and releases the new peptide.

| | Second Letter | | | | |
|---|--|--------------------------|--|--|------------------|
| First Letter | U | с | А | G | Third Letter |
| U | UUU UUC UUA UUA UUG | UCU UCC UCA UCG | UAU UAC UAA STOP ^b UAG STOP ^b | UGU UGC UGA STOP ^b UGG Trp | U C A G |
| с | CUU CUC CUA CUG | CCU CCC CCA CCG | $\left. \begin{matrix} CAU\\ CAC \end{matrix} \right\}_{His} \\ \left. \begin{matrix} CAA\\ CAG \end{matrix} \right\}_{Gln}$ | $\begin{pmatrix} CGU \\ CGC \\ CGA \\ CGG \end{pmatrix}$ Arg | U C A G |
| A | AUU AUC AUA AUG START ^a /Met | ACU ACC ACA ACG | $\left. \begin{array}{c} AAU\\ AAC \end{array} \right\} Asn \\ \left. \begin{array}{c} AAA\\ AAA \end{array} \right\} Lys \end{array}$ | $\left. \begin{array}{c} AGU \\ AGC \end{array} \right\} Ser \\ \left. \begin{array}{c} AGA \\ AGG \end{array} \right\} Arg \end{array}$ | U C A G |
| G | GUU GUC GUA GUG | GCU GCC GCA GCG | GAU GAC GAA GAG GAG | $\left. \begin{array}{c} GGU\\ GGC\\ GGA\\ GGG \end{array} \right\} Gly$ | U C A G |
| START? coden signals the initiation of a populate chain | | | | | |

START^a codon signals the initiation of a peptide chain STOP^b codons signal the end of a peptide chain.

| Site: Materials | Process | |
|--|--|--|
| Nucleus: nucleotides, RNA polymerase | A DNA template is used to produce mRNA. | |
| Cytoplasm: amino acids, tRNAs, aminoacyl–tRNA synthetase | Molecules of tRNA pick up specific amino acids according to their anticodons. | |
| Ribosome: Met–tRNA, mRNA, amino acyl–tRNAs | A start codon binds the first tRNA carrying amino acid methionine to the mRNA. Successive tRNAs bind to and detach from the ribosome as each amino acid adds to the protein. | |
| Ribosome: stop codon on mRNA | The protein is released from ribosome. | |
| | | |
| 0 | | |
| -GCG AGT GGA TAC- | | |
| | GC TCA CCT ATG— | |
| | | |
| | CG AGU GGA UAC— | |
| CGC UCA CCU AUG | | |
| —Ala—Ser—Gly—Tyr— | | |
| cytoplasmic pool of free tRNA, ready to be reactivated with a new amino acid | C G G U A U Ribosome | |
| | Nucleus: nucleotides, RNA polymerase Cytoplasm: amino acids, tRNAs, aminoacyl–tRNA synthetase Ribosome: Met–tRNA, mRNA, amino acyl–tRNAs Ribosome: stop codon on mRNA G G G G G G G G G G G G G G G G G G G | |

ribosome

advance

Try it.

 $C \rightarrow G$ $G \rightarrow C$

The following section of DNA is used to build mRNA for a protein:

-GAA - CCC - TTT -

A. What is the corresponding mRNA sequence? $A \rightarrow U$ $T \rightarrow A$

-CUU-GGG-AAA-

B. What are the anticodons on the tRNAs?

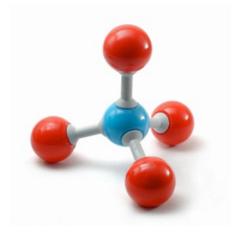
GAA for CUU CCC for GGG UUU for AAA

C. What is the amino acid order in the peptide?

-CUU-GGG-AAA-

| | Second Letter | | | | |
|--------------|---|--------------------------|---|--|------------------|
| First Letter | U | С | А | G | Third Lette |
| U | UUU UUC UUA UUA Leu | UCU UCC UCA UCG | UAU UAC UAA STOP ^b UAG STOP ^b | UGU UGC UGA STOP ^b UGG Trp | U C A G |
| с | CUU CUC CUA CUG | CCU CCC CCA CCG | $ \begin{array}{c} CAU \\ CAC \end{array} \hspace{-1.5mm} -1.5$ | $\left. \begin{array}{c} CGU\\ CGC\\ CGA\\ CGG \end{array} \right\}_{Arg}$ | U C A G |
| A | $\left. \begin{array}{c} AUU\\ AUC\\ AUA \end{array} \right\} IIe\\ AUG STARTa/Met \end{array} \right.$ | ACU ACC ACA ACG | $ \begin{array}{c} AAU \\ AAC \end{array} Asn \\ AAA \\ AAA \\ AAG \end{array} \right\} Lys $ | AGU AGC Ser AGA AGG Arg | U C A G |
| G | GUU GUC GUA GUG | GCU GCC GCA GCG | GAU GAC GAA GAG Glu | GGU GGC GGA GGG | U C A G |

START^a codon signals the initiation of a peptide chain. STOP^b codons signal the end of a peptide chain.



Ch17 Genetic Code Chain Elongation ► RNA Termination Differences from DNA Function & Structure **Mutation** Stems & Loops Mistakes happen Types of RNA Viruses Messenger (mRNA) Taking over the factory Transfer (tRNA) Ribosomal (rRNA) Protein Synthesis Genes are stored in DNA Transcription Transcription Code is written on mRNA Translation mRNA Codons / Genetic Code ► START & STOP signals Amino Acid Codes Protein Thr ▶ tRNA Anticodons Structure & Activation PPROV 40

Genetic Mutations

- If the enzyme that converts tyrosine to melanin is defective, no melanin is produced and a genetic disease known as albinism results.
- A peacock with albinism does not produce the melanin needed to make the bright colors of its feathers.



Mutations

- A mutation, or change in the nucleotide sequence of DNA, may
- alter the sequence of amino acids
- affect the structure and function of a protein in a cell
- Mutations may result from
- X-rays
- overexposure to sun (ultraviolet, or UV, light)
- chemicals called mutagens
- some viruses



Mutations

- If a mutation occurs in a somatic cell, a cell other than a reproductive cell, the altered DNA will be limited to that cell and its daughter cells.
- If the mutation causes uncontrolled growth, cancer could result.
- If a mutation occurs in a germ cell (egg or sperm), then all the DNA produced in a new individual will contain the same genetic change.



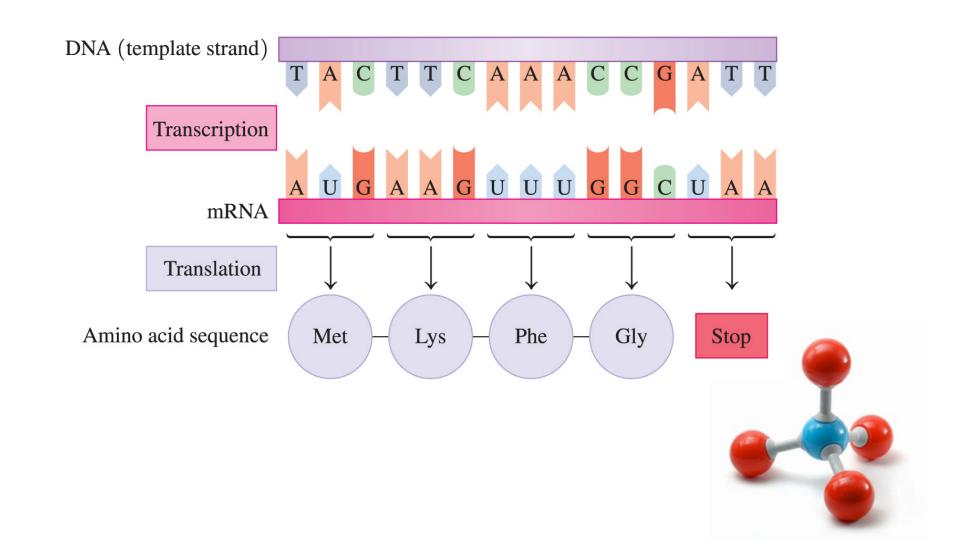
Types of Mutations

- A substitution or point mutation is the replacement of one base in the template strand of DNA with another.
- If a substitution or point mutation changes the nucleotide, a different amino acid may be inserted into the polypeptide.
- If this produces no change in the amino acid sequence, it is called a silent mutation.
- A frameshift mutation is the insertion or deletion of a single nucleotide into the sequence resulting in a change to all subsequent codons, leading to a new amino acid sequence.



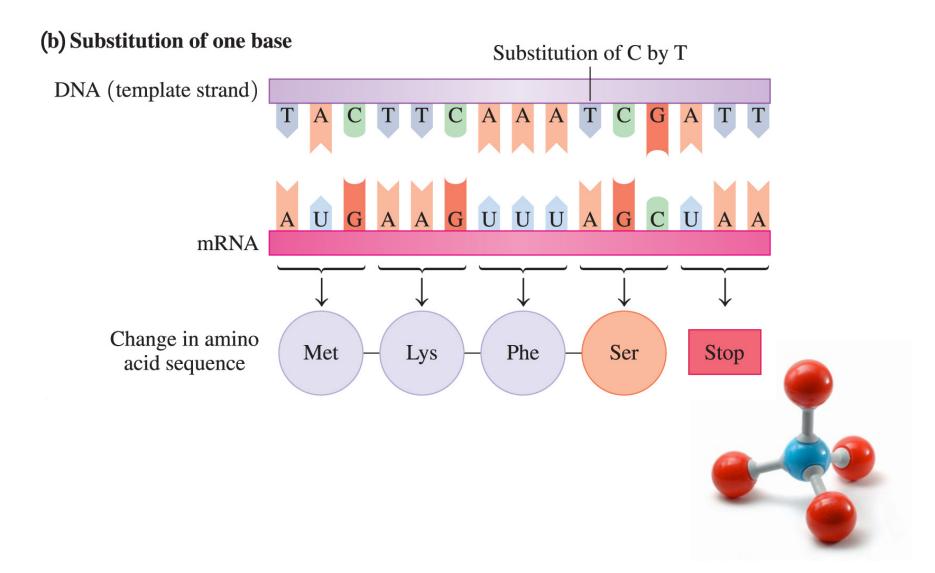
Normal DNA and Protein Synthesis

The normal DNA sequence produces an mRNA that provides instructions for the correct series of amino acids in a protein.



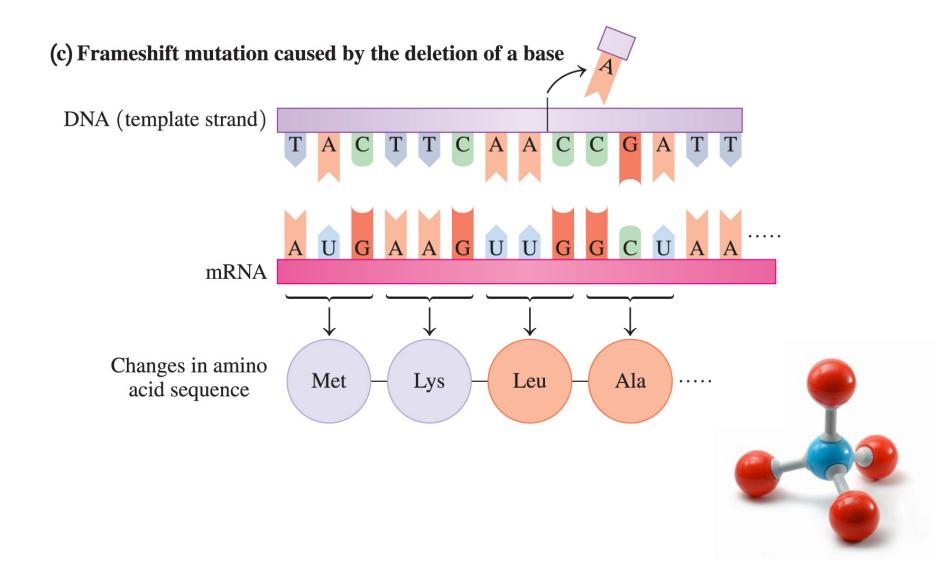
Mutation: Substitution

Substitution of a base in DNA changes a codon in the mRNA, leading to the placement of an incorrect amino acid in the polypeptide.



Frameshift Mutation

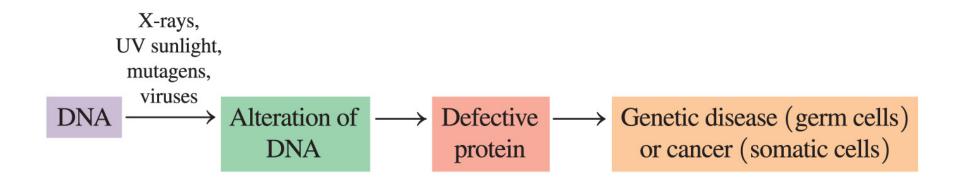
The deletion of a base causes a frameshift mutation, which changes the mRNA codons that follow the mutation and produces a different amino acid sequence.



Effect of Mutations

When a mutation causes a change in the amino acid sequence, the structure of the resulting protein may be severely altered, causing loss of its biological activity.

When this condition is hereditary, it is called a genetic disease.

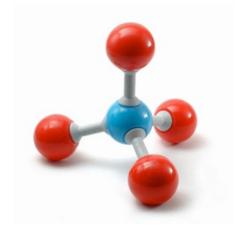




Genetic Diseases

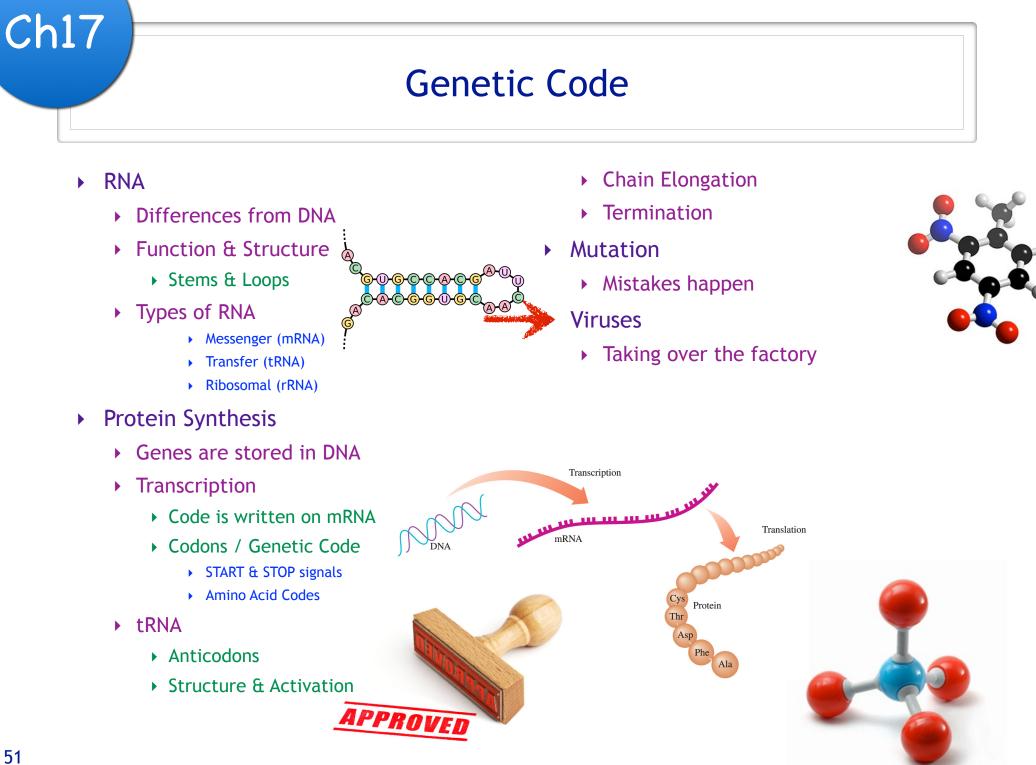
A genetic disease is the result of a defective enzyme caused by a mutation in its genetic code. For example,

- phenylketonuria (PKU) results when DNA cannot direct the synthesis of the enzyme phenylalanine hydroxylase, required for the conversion of phenylalanine to tyrosine
- albinism results when the enzyme that converts tyrosine to melanin is defective



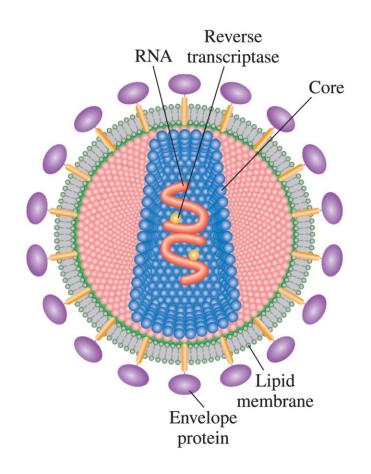
Genetic Diseases

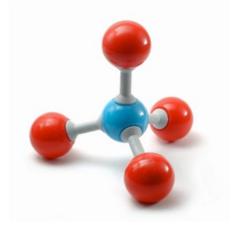
| Genetic Disease | Result |
|---------------------------------------|--|
| Galactosemia | In galactosemia, the transferase enzyme required for the metabolism of galactose-1-phosphate is absent, resulting in the accumulation of galactose-1-phosphate, which leads to cataracts and mental retardation. Galactosemia occurs in about 1 in every 50 000 births. |
| Cystic fibrosis (CF) | Cystic fibrosis is caused by a mutation in the gene for the protein that regulates the production of stomach fluids and mucus. CF is one of the most common inherited diseases in children, in which thick mucus secretions make breathing difficult and block pancreatic function. |
| Down syndrome | Down syndrome is the leading cause of mental retardation, occurring in about 1 of every 800 live births; the mother's age strongly influences its occurrence. Mental and physical problems, including heart and eye defects, are the result of the formation of three chromosomes (trisomy), usually number 21, instead of a pair. |
| Familial hypercholesterolemia | Familial hypercholesterolemia occurs when there is a mutation of a gene on chromosome 19, which produces high cholesterol levels that lead to early coronary heart disease in people 30 to 40 years old. |
| Muscular dystrophy (MD) (Duchenne) | Muscular dystrophy, Duchenne form, is caused by a mutation in the X chromosome. This muscle- destroying disease appears at about age 5, with death by age 20, and occurs in about 1 of 10 000 males. |
| Huntington's disease (HD) | Huntington's disease affects the nervous system, leading to total physical impairment. It is the result of a mutation in a gene on chromosome 4, which can now be mapped to test people in families with a history of HD. There are about 30 000 people with Huntington's disease in the United States. |
| Sickle-cell anemia | Sickle-cell anemia is caused by a defective form of hemoglobin resulting from a mutation in a gene on chromosome 11. It decreases the oxygen-carrying ability of red blood cells, which take on a sickled shape, causing anemia and plugged capillaries from red blood cell aggregation. In the United States, about 72 000 people are affected by sickle-cell anemia. |
| Hemophilia | Hemophilia is the result of one or more defective blood-clotting factors that lead to poor coagulation, excessive bleeding, and internal hemorrhages. There are about 20 000 hemophilia patients in the United States. |
| Tay–Sachs disease | Tay–Sachs disease is the result of a defective hexosaminidase A, which causes an accumulation of gangliosides and leads to mental retardation, loss of motor control, and early death. |



Viruses

• HIV causes AIDS, which destroys the immune system in the body. HIV is a retrovirus that infects and destroys T4 lymphocyte cells, which are involved in the immune response.

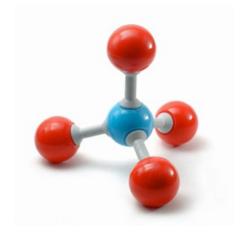




Viruses

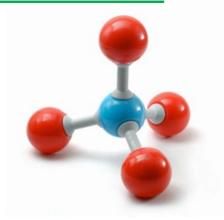
Viruses

- are small particles of 3 to 200 genes that require a host cell to replicate
- contain a nucleic acid-DNA or RNA, but not both-inside a protein coat
- do not have the necessary material, such as nucleotides and enzymes, to make proteins and grow
- replicate by invading a host cell and taking over the machinery and materials necessary for protein synthesis and growth



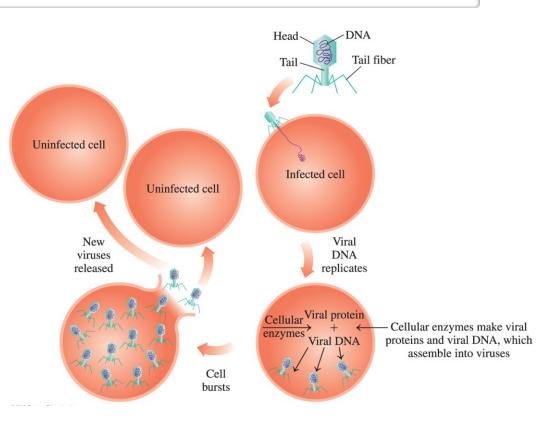
Some Diseases Caused by Viruses

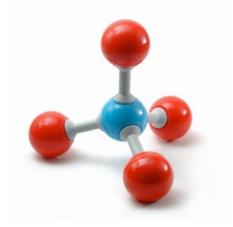
| Disease | Virus |
|-------------------------|---|
| Common cold | Coronavirus (over 100 types), rhinovirus (over 110 types) |
| Influenza | Orthomyxovirus |
| Warts | Papovavirus |
| Herpes | Herpesvirus |
| HPV | Human papilloma virus |
| Leukemia, cancers, AIDS | Retrovirus |
| Hepatitis | Hepatitis A virus (HAV), hepatitis B virus (HBV), hepatitis C virus (HCV) |
| Mumps | Paramyxovirus |
| Epstein–Barr | Epstein–Barr virus (EBV) |
| Chicken pox (shingles) | Varicella zoster virus (VZV) |



Viruses

- ► A virus
 - attaches to the host cell and injects its viral DNA
 - uses the host cell's amino acids to synthesize viral protein
 - uses the host cell's contents to make viral RNA
- When the cell bursts, the new viruses are released to infect other cells.

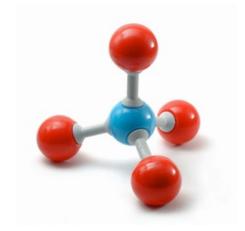




Viral Infection

A viral infection begins when

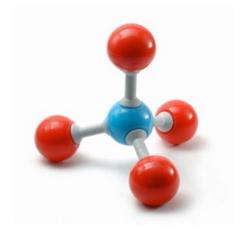
- a virus enzyme makes a hole in the host cell wall
- the viral nucleic acids enter and mix with the materials in the host cell
- the host cell begins to replicate the viral DNA (if the virus contains DNA)
- viral DNA produces viral RNA, and a protease processes proteins to produce a protein coat to form a viral particle that leaves the cell
- the cell synthesizes so many virus particles that it releases new viruses to infect more cells



Reverse Transcription

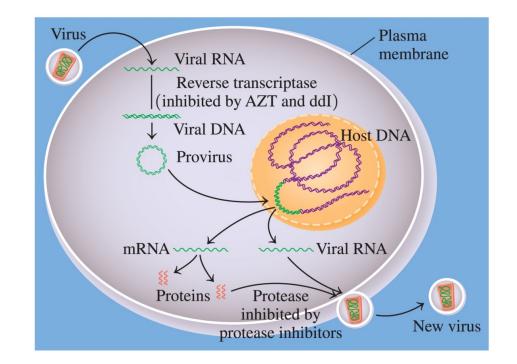
In reverse transcription,

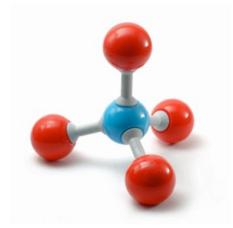
- a retrovirus, which contains viral RNA but no viral DNA, enters a cell
- the viral RNA uses *reverse transcriptase* to produce a viral DNA strand
- the viral DNA strand forms a complementary DNA strand
- the new DNA uses the nucleotides and enzymes in the host cell to synthesize new virus particles

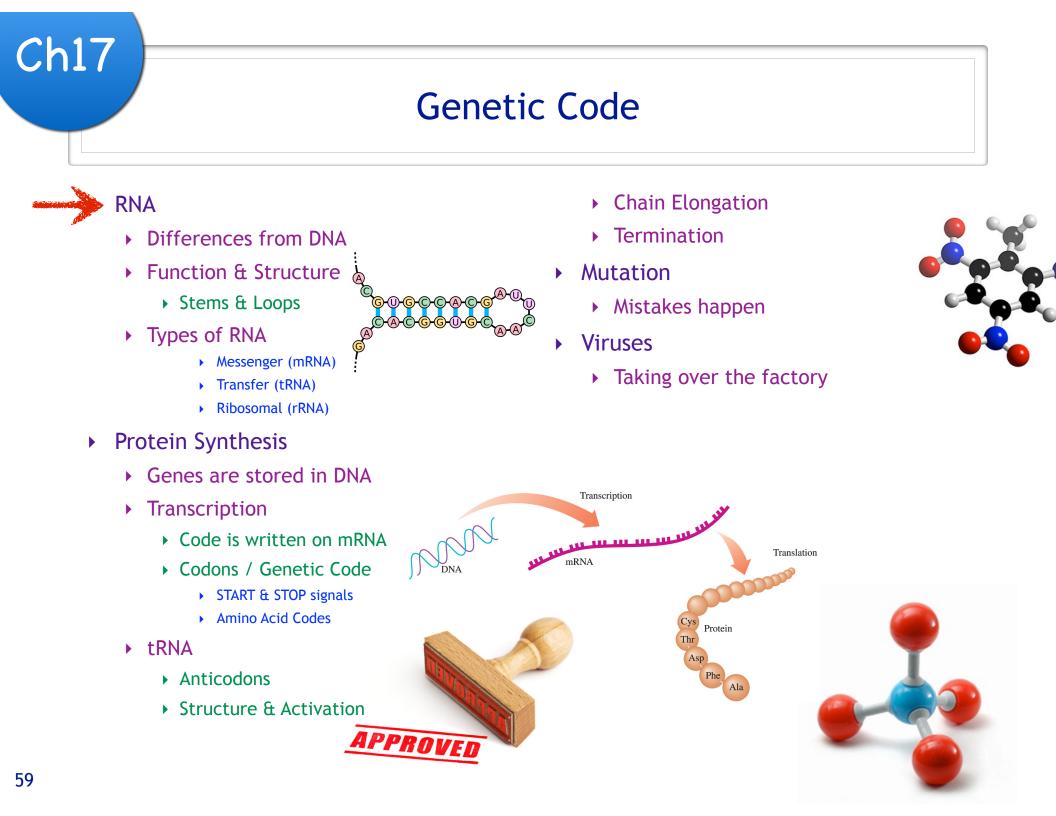


Reverse Transcription

- After a retrovirus injects its viral RNA into a cell, it forms a DNA strand by reverse transcription.
- The DNA forms a double-stranded DNA called a provirus, which joins the host cell's DNA.
- When the cell replicates, the provirus produces the viral RNA needed to produce more virus particles.







Questions?

