THURSDAY 11/29



REMINDERS

Next week

- -Monday December 3rd Unit 5 Vocabulary Quiz
- -Monday December 3rd AFTER SCHOOL Unit 4 Makeups (Quiz, Exam, Practical)
- -Tuesday December 4th Unit 5 Exam
- Wednesday December 5th Prefix/Suffix Quiz #6
- Thursday December 6th Unit 5 Practical Test

P.O.D.

- What are the 4 criteria that Watson & Crick knew they had to work with when developing their model of the structure of DNA?
- Identify (A), (B), (C), (D), (E), & (F)





The Four Criteria

- 1) The molecule must be able to store A LOT of information
- 2) The molecule must be able to copy itself with great precision
- 3) The molecule must be able to make mistakes when being copied
- 4) The molecule must be readable by cells





Remember that DNA is a double helix where one side of the "ladder" is the complement of the other side

- This is what makes precise replication possible



DNA helicase breaks the hydrogen bonds between base pairs, which "unwinds" and "unzips" the DNA





Each strand in the double helix acts as a template for the synthesis of a new and complementary strand





DNA polymerase adds complementary nucleotides one by one to both templates





This results in two "daughter" identical strands of DNA

- Each with one new strand and one old strand
 - Semiconservative







- Sometimes, a mistake is made when adding complementary base pairs and the wrong bases are paired together
 - That means that the mistake has the potential to change many things



Genes code for certain proteins and traits and can turn other genes off or on

- Mistakes can turn out to be good and they can even lead to no actual change, but they can also be deadly



About 1 in 10⁵ base pairs can result in an incorrect pairing

- DNA polymerase proofreads as it adds nucleotides
 - It can remove the vast majority of the mistakes that are made



What a mistake is good, neutral, or bad depends on how the DNA is read by the cell

- This is the next part we will talk about in a bit

RNA – another nucleic acid

- Ribonucleic acid
 - Very similar to DNA but...
 OHas ribose instead of deoxyribose
 - Single-stranded instead of double stranded
 - OUracil (pyrimidine, 1 ring of carbon, 2 hydrogen bonds) instead of thymine



RNA – another nucleic acid

RNA is the way that information is sent from the nucleus where DNA is stored to ribosomes where proteins are synthesized





Three types of RNA

mRNA

- messenger RNA: the RNA molecule transcribed from the DNA template
- tRNA
 - transfer RNA: RNA molecules that brings and binds amino acids to the ribosome and mRNA

rRNA

- ribosomal RNA: make up the ribosome (ribosomes are made out of RNA!)

Reading RNA

How many 3-letter 'words' can you make with the following 4 letters (they don't have to make sense)?

A

C G (you can use letters more than once)

A U C G

- AAA
- AUC
- AUG
- AUA
- ACG
- ACA
- AUA
- etc.

- UUU CCC
- UUA CGC GCG
- UUC
- UUG
- UAU
- UCU
- UGU
- etc.

- CUA
- CAU GCA

- GGG

- GCU

- CAC CUA
- CAG GUC
- CUA GAG
- etc. etc.

P.O.D			
	A T	C G	
- AAA	- TTT	- CCC	- GGG
- ATC	- TTA	- CGC	- GCG
AND EVEN MORE!			
- ALG	- IAI	- LAL	- CTA
- ACA	- TCT	- CAG	- GTC
- ATA	- TGT	- CTA	- GAG
- etc.	- etc.	- etc.	- etc.



- There are 4 letters and you want to make 3-letter words
- You may use letters multiple times
- To figure out how many possibilities there are....
 - Write out placeholders for how many letters you want in your word



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 - Write out placeholders for how many letters you want in your word
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 - Then, multiply those numbers together

4 x 4 x 4



- There are 4 letters and you want to make 3-letter words
- You may use letters multiple times
- To figure out how many possibilities there are....
 - Write out placeholders for how many letters you want in your word
 - Then write the number of letters that could be used in that spot
 - Then, multiply those numbers together

4 x 4 x 4 = 64



Reading DNA

Cells read DNA in groups of 3 base pairs at a time

- Each set of 3 base pairs is called a codon

•Codons code for amino acids

- One amino acid can be coded for by multiple codons
 - ... but each single codon will always code for the same amino acid







mRNA message transcribed from DNA: UUUUACAGGGCCGUAGAAUGA



Ribosome reads the mRNA message: UUU UAC AGG GCC GUA GAA UGA



Ribosome reads the mRNA message: Phy UAC AGG GCC GUA GAA UGA



Ribosome reads the mRNA message: Phy | Tyr | AGG | GCC | GUA | GAA | UGA



Ribosome reads the mRNA message: Phy | Tyr | Arg | GCC | GUA | GAA | UGA



Ribosome reads the mRNA message: Phy | Tyr | Arg | Ala | GUA | GAA | UGA



Ribosome reads the mRNA message: Phy Tyr Arg Ala Val GAA UGA



Ribosome reads the mRNA message: Phy Tyr Arg Ala Val Glu UGA


Ribosome translates to amino acids: Phy | Tyr | Arg | Ala | Val | Glu | STOP



The protein chain:



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Use the Codon to Amino Acid chart to translate the following mRNA message...

AUGAAACGAGGGCUGUUUCGAAAAUGA



AUG AAA CGA GGG CUG UUU CGA AAU UGA



Met AAA CGA GGG CUG UUU CGA AAU UGA



Met Lys CGA GGG CUG UUU CGA AAU UGA



Met Lys Arg GGG CUG UUU CGA AAU UGA



Met | Lys | Arg | Gly | CUG | UUU | CGA | AAU | UGA



Met | Lys | Arg | Gly | Leu | UUU | CGA | AAU | UGA



Met | Lys | Arg | Gly | Leu | Phy | CGA | AAU | UGA



Met | Lys | Arg | Gly | Leu | Phy | Arg | AAU | UGA



Met Lys Arg Gly Leu Phy Arg Asp UGA



Met | Lys | Arg | Gly | Leu | Phy | Arg | Asp | STOP





Met Lys Arg Gly Leu Phy Arg Asp STOP

Met Lys Arg Gly Leu Phy Arg Asp

The START codon corresponds with Methionine But

The STOP codon does not correspond with an amino acid



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Transcription

- The first step of cells 'reading' the instructions in DNA and carrying them out
 - The goal is to make an RNA copy of a gene's DNA sequence

 Cells do transcription with single genes and groups of genes, not with an organisms' whole genome



RNA polymerase binds to the START codon at the beginning of the gene and then separates the DNA strands and exposes a single-strand of DNA that will serve as a template







RNA polymerase "reads" the template strand of the gene one base at a time and then builds an RNA copy by adding complementary bases

DNA rewind

RNA transcript

DNA unwind

b) Elongation





RNA polymerase "reads" the STOP codon and the RNA transcript is complete

This is called mRNA









This strand of mRNA leaves the nucleus and carries its message to the ribosome ribosomes mRNA nascent proteins 100 nm



- After the mRNA is complete, it travels to the ribosomes where the mRNA is read codon by codon
 - As each codon is read, the corresponding amino acid is bonded









A chain of amino acids forms the peptide (protein) that the gene that was transcribed/translated coded for





A ribosome assembles around the mRNA and a tRNA carrying the START amino acid





Then, the ribosome reads the codons one at a time (3 base pairs at a time) and tRNA brings in the corresponding amino acids, adding them to the protein chain





When the ribosome reads a STOP codon, the protein chain separates from the mRNA and ribosome







After a newly synthesized protein chain gets released, it may still need to

- Fold into the right shape
- Undergo processing
- Get shipped to the correct place in the cell
- Combine with other protein chains

How to think about all this....



Watson & Crick's 4 Criteria



Criteria 1

The molecule must be able to store A LOT of information

How does DNA store information?



Criteria 2

The molecule must be able to copy itself with great precision

Why is this important?How is it done?



Criteria 3

The molecule must be able to make mistakes when being copied

Why are mistakes important?
How can DNA make mistakes?


Criteria 4

The molecule must be readable by cells

How does the cell "read" the DNA?

- What does it do with the information?
- What is transcription?
- What is translation?

Transcription **R Translation** Practice



Steps:

- (1) Transcribe DNA using mRNA
 - Remember: RNA has U and no T
- (2) Group the mRNA into codons
 - Groups of 3
- (3) Translate mRNA into amino acids
 - Use the chart
- (4) String the amino acids together in to a protein chain
 - The START codon corresponds to an amino acid but the STOP codon does not

DNA sequence – a gene ATGTCTACTAATGGGAGTTACTTAGAGTAG



mRNA – transcription of DNA AUGUCUACUAAUGGGAGUUACUUAGAGUAG















Ribosome translates codon to amino acids Met | Ser | Thr | Asp | Gly | Ser | UAC | UUA | GAG | UAG



Ribosome translates codon to amino acids Met | Ser | Thr | Asp | Gly | Ser | Tyr | UUA | GAG | UAG



Ribosome translates codon to amino acids Met | Ser | Thr | Asp | Gly | Ser | Tyr | Leu | GAG | UAG



Ribosome translates codon to amino acids Met | Ser | Thr | Asp | Gly | Ser | Tyr | Leu | Glu | UAG



Ribosome translates codon to amino acids Met | Ser | Thr | Asp | Gly | Ser | Tyr | Leu | Glu | STOP





String amino acids together into a protein chain

Met | Ser | Thr | Asp | Gly | Ser | Tyr | Leu | Glu | STOP





Transcribe and Translate the following messages:

- I. ATGGTTAATCCCCACCGATTTTGTTGAAGTGA
- 2. ATGCATGAGGGGTTTCTTATTGTGAATAAATAG
- 3. ATGTATCGCAGTGATGTTCTTCCAGGCGGGTAA

Work your way back to the original DNA message

- 1. Met-Thr-Ala-Tyr-Asp-Ser-Gly-Iso-Leu-Val-STOP
- 2. Met-Pro-Ser-Cys-Gly-Asp-Ala-Phy-Ser-Pro-STOP
- 3. Met-Arg-Cys-Ser-Pro-Thr-Gly-Iso-Leu-Glu-STOP