Name	Class	Date

Skills Worksheet

Protein Synthesis

INTERPRETING TABLES

Use the table below to complete items 1-17.

	Codons in mRNA						
First base	Second base U C A G			Third base			
U	UUU UUC Phenylalanine UUA Leucine	UCU UCC UCA UCG Serine	UAU Tyrosine UAA UAA Stop	UGU UGC Cysteine UGA – Stop UGG – Tryptophan	U C A G		
С	CUU CUC CUA CUG	CCU CCC CCA CCG	CAU Histidine CAC Glutamine CAA Glutamine	CGU CGC CGA CGG	U C A G		
A	AUU AUC AUA Isoleucine AUA - Start	ACU ACC ACA ACG Threonine	AAU Asparagine AAA AAA AAG Lysine	AGU Serine AGA AGG Arginine	U C A G		
G	GUU GUC GUA GUG	GCU GCC GCA GCG	GAU Aspartic GAC acid GAA Glutamic GAG acid	GGU GGC GGA GGG	U C A G		

Complete the table below showing sequences of DNA, mRNA codons, anticodons, and corresponding amino acids. Use the list of mRNA codons in the table above to assist you in completing this exercise. Remember that the genetic code is based on mRNA codons.

Decoding DNA				
DNA	1	2	GAT	3
mRNA codon	4	5	6	UAU
Anticodon	7	UUC	8	9
Amino acid	Tryptophan	10	11	12

Name	Class	Date
Protein Synthesis contin	ued	
the mRNA codons in the t below. In the space provide	able on the previous led, write the names	ct each amino acid sequence. Use spage to complete items 13–16 of the amino acids that correspond An example is provided for you.
Example:		
mRNA sequence:	UGU-CCG	cysteine-proline
mutation sequence:	UGC-CGC	cysteine-arginine
13. mRNA sequence:	GAA-CGU	
mutation sequence:	GAU-CGU	
14. mRNA sequence:	AUC-UGC	
mutation sequence:	AUC-UGG	
15. mRNA sequence:	UGU-CCU-CCU	
mutation sequence:	UGU-UUC-CCU	
16. mRNA sequence:	GGG-UUA-ACC	
mutation sequence:	GGU-UAA	
17. What kind of mutation Explain.	n occurred to the mI	RNA sequence in item 16 above?

DNA Structure

INTERPRETING DIAGRAMS

- 1. A. phosphate group
 - **B.** pyrimidine (thymine)
 - C. hydrogen bond
 - **D.** purine (adenine)
 - E. deoxyribose
- 2. the hydrogen bonds between the bases; cytosine and guanine form three hydrogen bonds, whereas adenine and thymine form only two hydrogen bonds.
- 3. TAA-GGC

Protein Synthesis

INTERPRETING TABLES

- 1. ACC
- **2.** TTC
- **3.** ATA
- 4. UGG
- **5.** AAG
- **6.** CUA
- **7.** ACC
- **8.** GAU
- **9.** AUA
- **10.** lysine
- 11. leucine
- 12. tyrosine
- **13.** glutamic acid-arginine to aspartic acid-arginine
- **14.** isoleucine-cysteine to isoleucine-tryptophan
- **15.** cysteine-proline-proline to cysteine-phenylalanine-proline
- **16.** glycine-leucine-threonine to glycine-stop
- 17. A frameshift mutation occurred.

 One of the G nucleotides and two C nucleotides have been deleted. The second codon is a stop codon, which will cause translation to end prematurely. The protein for that gene will be shortened and incomplete.

Gene Technology

INTERPRETING DIAGRAMS

- 1. The structure labeled *A* is a plasmid. It is removed from the bacterial cell so that the plasmid can be used as a vector to carry the insulin gene into a bacterial cell.
- 2. Restriction enzymes are used to cut DNA. DNA molecules cut with restriction enzymes have sticky ends that allow different DNA fragments cut with the same restriction enzyme to combine.
- **3.** This is recombinant DNA. (The DNA with the insulin gene and plasmid DNA are combined).
- **4.** This is a bacterial cell that contains recombined plasmids (plasmids containing the insulin gene).
- 5. CCGG and GGCC
- **6.** pair on the left—GGCC/CCGG; pair on the right—CCGG/GGCC
- 7. Tetracycline, an antibiotic, destroys bacterial cells. Some bacterial cells, however, contain a gene for tetracycline resistance in their plasmid DNA, and they are not harmed by the antibiotic. These cells are called tetracycline-resistant cells. In the diagram, the plasmid DNA used in the genetic engineering experiment has the gene for tetracycline resistance. Only the cells that have taken up the plasmid DNA with the gene for tetracycline resistance survive when tetracycline is added to the cultures. Only the resistant cells, those that also carry the gene of interest, survive.