Name		Period Date									
	tion, Transcription and Translati										
Review question											
 Transcription and t 	ranslation is also known as prote	ein synthesis, and is the expression of genes . The geneti	c code								
determines the amino acid sequence of a polypeptide, and the properties of the amino acids give the final structure											
and function of the	protein. Other than membrane	proteins, state four functions of proteins in the cell.									
2.7.U4 Transcription	is the synthesis of mRNA conject	from the DNA base sequences by RNA polymerase.									
		including the roles of RNA polymerase, ribonucleoside									
•	complementary base pairing.	p.,									
	the synthesis of polypeptides or										
3. Complete the table	to compare and contrast the pr	ocesses of transcription and translation.									
	Transcription	Translation									
Begins with		mRNA									
Ends with											
Location											
Uses	RNA polymerase										
0383	NIVA polymerase										
	·	the process of translation. Outline the structure of the ri	ibosome								
and explain now it	is adapted to carry out translation	on.									
2.7.U6 The amino ac	id sequence of polypeptides is d	etermined by mRNA according to the genetic code.									
2.7.U7 Codons of thr	ee bases on mRNA correspond t	o one amino acid in a polypeptide.									
5. Define mRNA in ter	rms of it's function										
6. Suggest why the le	ngth of mRNA molecules varies.										
7. Describe what is m	eant by the term 'genetic code'.										
8. Define the term co											
	per of different codons combinat										
	of amino acids that can be transl										
11. Explain what is mea	ant by the term degenerate. Ref	er to the last two questions in your answer.									
2.7.U8 Translation de	epends on complementary base	pairing between codons on mRNA and anticodons on tR	RNA.								
12. State the molecule	on which anti-codons, which are	e complementary to codons, can be found.									
13. Complete the steps	s to outline the process of transl	ation.									
a. mRNA binds to	the of the r	ibosome.									
b. The mRNA con	tains a series of	each of which codes for an amino acid.									
		re complementary to the on the									
d. tRNA molecule	s bind to a	that corresponds to the anticodon									
e. The	binds to the small subunit	of the ribosome.									
		of the ribosome, but only can contain tR	RNA								
molecules at a		d procents codons in the first two									
g. The	moves along the mkNA an	d presents codons in the first two									

h.	with a	nticc	don \	s					_ to the codons bind (the bases are linked by the formation of														
i. j.		is formed between the two amino acids (carried by the tRNAs) moves along a tRNA moves to the binding site and																					
k. I.	Another The process (i.															nd _					_ is f	orm	ed.
2.7.S1	Use a table o	of the	gen	etic	cod	e to	dedı	uce v	which	ո cod	lon(s) cor	resp	ono	ls to	whic	:h an	nino	acid	<u>.</u>			
2.7.S3	Use a table o	of mF	RNA	codo	ns a	nd t	heir	corr	espo	ndin	g am	ino a	acid	s to	dedu	ıce tl	ne se	eque	ence	of an	nino	acid	S
coded l	oy a short mRN	A str	and	of kr	now	n bas	se se	que	nce.														
2.7.S4	Deducing the	e DN	A ba	se se	eque	nce	for t	he n	nRN <i>A</i>	\ stra	nd.												
	e genetic code to the questions			elp				Th	e ge	netic	cod	e – ł	now	mR	NA c	odoı	ns tra	ansl	ate t	o am	ino	acid	S
14. Dec	duce the codon	(s) th	nat ti	rans	late				http	. / /		h hic	nin	ia co	.m. a	/ N	10dis	. /	notic		la in	0.0	
for	Aspartate.								пцр	://w	VV VV . I		ond l		IIII.a	<u>u/_iv</u>	reur	a/ge	пец				
15. If m	nRNA contains	the b	ase	sequ	ienc	e			l	U		C	ona i	A		(3				Ke	у.	
CUGACUAGGUCCGGA										} Phe	UC			UAU)	Tvr	UGU	UGU Cys			Ala = Alanine (A) Arg = Arginine (R) Asn = Asparagine (N) Asp = Aspartate (D)			
a.	deduce the an	the amino acid sequence of									UC	UCC UCA UCG			TOP	UGA	C A G						
	the polypeptic	le tra	nsla	ted.					UUG Leu					UAG S	TOP	UGG				Cys = Cysteine (C) Gin = Glutamine (Q)			
b.	deduce the ba	se se	que	nce	of th	ie		С	CUC	CUC	CC	CCU CCC CCA CCG		CAU }	His	CGU CGC CGA CGG		C		Glu = Glutamate (E) Gly = Glycine (G) His = Histidine (H) Ile = Isoleucine (I)			
	DNA antisense	stra	nd f	rom	whi	ch	,		CUA					CAA }	GIn			G	Third				
	the mRNA was	s trar	nscril	bed.				List	AUU AUC AUA		AC			AAU]		AGU	U	btter	Leu = Leucine (L) Lys = Lysine (K)			(L)	
C.	If mRNA conta	ins t	he b	ase				A		J	ACC ACA				ASN AA Lys		AGC Ser AGA AGG Arg		۳	Met = Methionine (M) Phe = Phenylalanine (F Pro = Proline (P) Ser = Serine (S)			
	sequence ACU	AAC	ded	uce 1	the l	oase			AUG Met		AC	GCU GCC			Lys	AGG	Arg	G					
	sequence of th	ne Di	NA se	ense	stra	nd.			GUU)		Asp				GGT GGC		U		Thi	Thr = Threonine (T) Trp = Tryptophan (W)			
16. Tra	nscribe and tra	nslat	e th	is DN	۱A			G	GUA GUG	Val	GC	A Ala		GAA)	Glu	GGA GGG	Gly	A		Tyr	= Ty	rosine	(Y)
seq	Juence.																			Val	= Val	ine (V)
	DNA	Т	Α	С	G	G	G	С	С	С	G	T	G	Α	С	Α	G	С	С	Α	С	Т	
	mRNA																						
	Amino acid			<u> </u>		1			1							1			1				-
		<u> </u>						<u> </u>									<u> </u>			<u> </u>			J

- 17. An mRNA strand has 76 codons. How many amino acids will be in the polypeptide?
- 18. A polypeptide contains 103 amino acids. What is the length of the gene (unit = base pairs)?
- 19. A gene is 105kbp (kilobase pairs). How many amino acids are in the polypeptide?

2.7.A2 Production of human insulin in bacteria as an example of the universality of the genetic code allowing gene transfer between species.

20. Diabetes in some individuals is due to destruction of cells in the pancreas that secrete the hormone insulin. It can be treated by injecting insulin into the blood. Despite the differences in the amino acid sequence between animal and human insulin, they all bind to the human insulin receptor and cause lowering of blood glucose concentration. However, some diabetics develop an allergy to animal insulin, so it is preferable to use human insulin. In 1982 human insulin became commercially available for the first time. It was produced using genetically modified E. coli bacteria. Since then methods of production have been developed using yeast cells and more recently safflower plants. Describe what is meant by the term 'universality of the genetic code'.