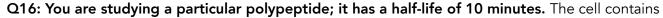
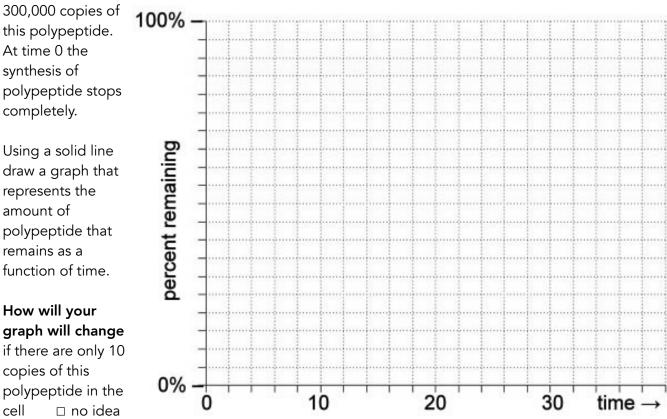
Q1: If genetic information were encoded in the		6
cells, rather than in the nucleotide sequences v Griffith's studies on the transformation in bacto		50
☐ A. would have produced exactly the same resu		9 (0
☐ B. would not have worked at all ☐ no i	,	
☐ C. would have identified proteins as the genetic		
Explain (below) why the <u>correct</u> answer is corre		
Explain (Bolon, III) the <u>solute</u> unoner to come		
Q2. In his studies, Griffith found that S-strain (s	mooth + virulent) bacteria	a grown in culture very
occasionally gave rise to R-strain (rough + aviru	•	rom $S \rightarrow R$ ).
Can you predict the relative frequency of a R -		
$\square$ A. The same as the S $\rightarrow$ R rate	☐ B. Much higher than th	
$\square$ C. Much lower than the S $\rightarrow$ R rate	□ D. impossible to say	□ no idea
Explain the logic of your answer		
Q3: A mutation occurs that leads to higher mu	tation rates in actively div	iding cells, but which has no
obvious effect on DNA in non-dividing cells. You	ou would be justified in assi	uming that the original
mutation inactivated		
$\ \square$ A. DNA-dependent DNA polymerase		
☐ B. DNA polymerase's proof-reading activity		
$\hfill \square$ C. DNA-dependent, RNA polymerase (primase	e)	
$\hfill\Box$ D. the repair of mutations due to the demethyl	ation of C's	□ no idea
Explain the logic behind your answer:		
Q4: PICK THE WRONG ANSWER: Which of the	following statements is c	orrect about DNA
replication?		n id
☐ A. DNA synthesis of the daughter strand alway		□ no idea
☐ B. DNA synthesis of the daughter strand alway	•	L'arta la consultante d
☐ C. DNA synthesis can occur in either direction	, ,	<b>'</b>
Explain the logic behind your answer (Hint: Dra	aw a picture with labels ar	id arrows indicating
synthesis directionality for full credit)		
Q5: The YUM gene is normally expressed only	in the skin cells of an orga	anism. In your studies, you
discover a mutant allele that leads to the expre	ession of the normal YUM	gene product in all cells of
the organism. Which is the most plausible exp		
$\ \square$ A. the mutation is in the regulatory region of the	=	□ no idea
$\hfill\Box$ B. the mutation is within the coding region of t	he YUM gene	
$\hfill \square$ C. the mutation alters DNA synthesis, leading t	to defect in primer synthesi	S
Explain the logic behind your answer		

@ biofundamentals project

Q6: As the percentage of GC in a double-stranded DNA completely and totally confident will occur?	A molecule increases, what would you be
☐ A. The rate of DNA synthesis will increase	□ no idea
☐ B. The mutation rate will increase	
$\hfill\Box$ C. The separation of two strands of the DNA molecule,	due to thermal motion, will increase
☐ D. The percentage of A in the DNA would decrease	
Explain the logic behind your answer	
Q7: A mutation occurs that leads to very high numbers of of a double-stranded DNA molecule, but with no obvious model for this effect would be to assume that the mutation □ A. the proof-reading activity associated with DNA polyr □ B. the DNA ligase	effects on the parental strands. A plausible n inactivated
☐ C. DNA-dependent, DNA polymerase	
☐ D. topoisomerase I	□ no idea
Explain the logic behind your answer:	
<ul> <li>Q8: Which is correct? the binding of a transcription fa</li> <li>□ A. has no effect on the direction of transcription</li> <li>□ B. determines exactly where translation begins</li> <li>□ C. determines where in the cell the encoded polypeptic</li> <li>□ D. determines which strand will be used to generate an</li> <li>□ E. determines when and where RNA primers are synthe</li> <li>Explain what will happen to the transcript (RNA) made if y reinsert back into to DNA the transcription factor's binding</li> </ul>	□ no idea  de will end up  RNA sized rou were able to remove, rotated 180°, and
Q9: Consider a cell. Which of the following processes a	re absolutely required to produce a
functional transcription factor?	
☐ A. DNA replication	□ no idea
☐ B. transcription	
☐ C. translation	
□ D. both transcription and translation	
<b>Explain</b> the logic of your answer.	
Q10: A protein has a short half-life, meaning that	
☐ A. it is rapidly synthesized	□ no idea
☐ B. it is rarely synthesized	
☐ C. the mRNA that directs its synthesis is unstable	
☐ D. it is rapidly degraded after it has been synthesized	
<b>Explain</b> the logic of your answer.	

Q11: You are asked to genetically engineer an organism so that it now amino acid (not one of the normally used set of amino acids). Which moved you NOT need to change?	
☐ A. one of the genes encoding a tRNA	□ no idea
☐ B. the genes that encode the ribosome	
☐ C. the gene that encodes the enzyme that adds the new amino acid to	the tRNA
☐ D. the genes encoding the enzymes involved in synthesizing the new a not normally made by the organism)	
<b>Explain</b> the logic of your answer.	
Q12: We discussed a type of mutation that allows a stop codon to be mutation would occur in a gene that encodes a	read as an amino acid. Such a
□ A. ribosomal RNA	□ no idea
☐ B. messenger RNA	
☐ C. transfer RNA	
☐ D. a gene's regulatory region	
<b>Explain</b> the logic of your answer (and why the other choices are wrong).	
Q13: The time between the synthesis and degradation of particular R (stochastic), like radioactive decay, because	NA or protein is noisy
☐ A.it depends upon random collisions between molecules	□ no idea
☐ B. it is determined by the molecule's structure	
☐ C. it is based on radioactive decay	
☐ D. it can be regulated by other factors	
<b>Explain</b> the logic of your answer.	
Q14: You isolate total <u>tRNA</u> from a cell and analyze its base composit nucleotides). This ratio will be	ion (i.e. the ratio of the various
□ A. A = U	□ no idea
□ B. A = G	
☐ C. the same as the bulk composition of the cell's DNA (but with Us inst	tead of Ts)
☐ D. impossible to know based on the information supplied	
<b>Explain</b> the logic of your answer.	
Q15: A mis-sense mutation can alter a polypeptide's 3D folding beca	use
☐ A. a different amino acid is inserted at the site of the mutation	□ no idea
☐ B. the polypeptide's synthesis stops prematurely	
☐ C. any change at any position of a polypeptide will lead to misfolding	
☐ D. it will alter the rate at which mRNA is synthesized	
<b>Explain</b> the logic of your answer.	





How might a cell could benefit from making a protein with a short half life?

## Q17: For an organism to be able to survive a mutation that creates a non-sense suppressor, which must be true?

be tide:	
□ A. the mutated gene must be relatively unimportant	□ no idea
$\square$ B. the original mutation (the mutation that is suppressed) must be in a non-coding re	gion
□ C. there must be multiple genes encoding specific tRNAs	
$\hfill \square$ D. the mutation must alter the region of the tRNA that determines which amino acid	is attached to the
tRNA	

**Explain** the logic of your answer.

## Q18: A non-sense mutation will always ...

- ☐ A. lead to the production of a longer polypeptide ☐ no idea ☐ B. lead to the production of a shorter polypeptide
- ☐ C. lead to the production of a dysfunctional polypeptide
- □ D. generally have no effect on polypeptide function

**Explain** how the position of a non-sense mutation would be likely to influence polypeptide activity.

Q19: You are studying the XUP gene of the speckled trout (a euka negatively acting transcription factor. You identify a mutation in the X mutant Xup protein is secreted from the cell. Which is the most likely whose transcription is directly regulated by the Xup protein?	UP gene and you find that the
☐ A. no effect, since it normally acts negatively	□ no idea
☐ B. their expression would increase	
☐ C. their expression would decrease	
$\hfill\Box$ D. the expression of all genes would increase	
<b>Explain</b> the logic of your answer.	
Q20: A polypeptide passes through a membrane once, and only of the case for the region of the polypeptide that that passes through $\square$ A. an unstructured polypeptide with both hydrophobic and hydroping B. a $\beta$ -sheet like structure with hydrophilic R-groups $\square$ C. an $\alpha$ -helix with hydrophobic R groups $\square$ D. it is impossible to make a plausible model based on the information.	h the membrane? In will form hilic R-groups □ no idea
<b>Explain</b> the logic of your answer.	
Q21: Draw and label: How could a membrane channel protein be polypeptide?	generated using a single □ no idea
Q22: If you were trying to devise a simple system that could recognize proteins in the cytoplasm of a cell, you might look for	gnize unfolded (denatured)
☐ A. acidic amino acid residues within the interior of the molecule	□ no idea
☐ B. non-peptide bonds on the surface	
□ C. a net positive surface charge	
D. multiple hydrophobic R groups on the surface  Total in the last of the surface of the surfac	
<b>Explain</b> the logic of your answer and describe what is likely to happer refolded correctly.	n it the untolded protein is not
	-:
Q23: A protein kinase phosphorylates a normally cytoplasmic prot the protein is found in the nucleus. Which of the following most li	
•	no idea
□ B. phosphorylation activates a signal sequence	
☐ C. phosphorylation activates a nuclear localization sequence	
☐ D. phosphorylation activates a nuclear export sequence	
<b>Explain</b> the logic of your answer.	

Expression of the ZIP gene 1		nt gene	r is regu	nated by	y the tra	mscrip	uon ia	ctor z	.IF.
depends on another	4								
transcription factor, ZNG.									
The ZNG gene is always expressed, but the ZNG protein is active only when the allosteric effector molecule ZOUP is present.	percent accumulating								
At time = 0 we add enough ZOUP to the culture to activate all of the ZNG protein present. Draw a graph of the accumulation of the CHL polypeptide as a function of									
time after the addition of ZOUP.	0% –		<del>-i-i-</del>	<del>       </del>	<del>-i i i</del>	<del>- i- i-</del>	<del>-i i</del>	time	iii
□ no idea  Describe the assumptions you m  Q25: Some genes are transcribed and translated.					of RNA	that is	both	transc	ribed
<ul><li>□ A. mRNAs</li><li>□ B. rRNAs</li><li>□ C. tRNAs</li><li>□ D. depends on the gene</li></ul>						□ nc	idea		
<b>Explain</b> the logic of your answer (i	nclude why	are the v	vrong ch	noices w	rong).				
Q26: How is regulation by an allo	osteric effe	ctor diff	erent fro	om regu	llation b	y proto	eolytic	cleav	age?
<ul><li>□ A. irreversible</li><li>□ B. reversible</li><li>□ C. always positive</li><li>□ D. always negative</li></ul>			1	□ no ide	ea				
<b>Explain</b> the logic of your answer.									

Q27: A mutation occurs that replaces an mRNA's normal start codon with a stop codon. Draw and explain what can happen ....