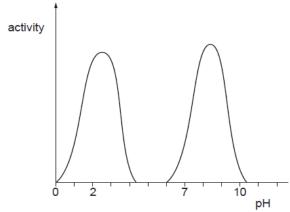
organisms. Enzymes behave a catalyse only one particular re-	
	ork better on heating, but enzymes rarely work at C. Explain why this is the case.
(b) Using the shape below to r	[2] represent an enzyme, sketch how an enzyme is specific to the
breakdown of a particular subs	
enzyme + substrate (c) Describe the effects of a co	enzyme-substrate complex enzyme + products ompetitive, and of a non-competitive inhibitor on the interaction
between enzyme and substrat	·
(d) (i) The diagram shown illus	[2] strates an enzyme-catalysed reaction. On the diagram sketch
the graph that would be obtain	ed if the same reaction was carried out in the presence of a
non-competitive inhibitor.	
initial reaction	
rate/mol dm ⁻³ s ⁻¹	
Į.	/concentration of substrate / mol dm ⁻³

(ii) Explain why a non-competitive inhibitor has this effect on the reaction.
(June 2011 P41 Q7)
Q2 Enzymes are protein molecules that are highly efficient in catalysing specific chemical reactions in living organisms. (a) To work in tissues, enzyme molecules generally need to be water-soluble. What does this tell you about the nature of the side-chains on the exterior of the molecules?
(b) Enzymes function by a substrate molecule interacting with a particular part of the enzyme known as the 'active site'. The substrate is converted into products that are then released, to be replaced by another substrate molecule. (i) Describe briefly the primary, secondary and tertiary structures of an enzyme.
(ii) The activity of an enzyme depends upon the tertiary structure of the protein molecule. Explain how the tertiary structure produces an effective active site.
(iii) Give two conditions that can reduce the activity of an enzyme, explaining the reason in each case.
1
II
(c) An individual enzyme operates best at a specific pH. Different enzymes operate best under conditions of different pH. Three enzymes involved in the digestion of food are amylase, pepsin and trypsin.

- Amylase, found in saliva, hydrolyses starch to a mixture of glucose and maltose under approximately neutral conditions.
- Pepsin hydrolyses proteins to peptides in the acid conditions of the stomach.
- Trypsin continues the hydrolysis of peptides to amino acids in the mildly alkaline conditions of the small intestine.

The graph below shows the activity of two of the three enzymes mentioned above.



- (i) Label each peak shown with the name of the enzyme responsible, either amylase, pepsin or trypsin.
- (ii) On the axes above, sketch the graph that the third enzyme would produce, and label it with the name of that enzyme.

(June 2011 P42 Q6)

- Q3 Proteins exist in an enormous variety of sizes and structures in living organisms. They have a wide range of functions which are dependent upon their structures. The structure and properties of an individual protein are a result of the primary structure – the sequence of amino acids that form the protein.
- (a) Proteins are described as condensation polymers.
- (i) Write a balanced equation for the condensation reaction between two glycine molecules, H2NCH2CO2H.
- (ii) Draw the skeletal formula for the organic product.
- **(b)** X-ray analysis has shown that in many proteins there are regions with a regular arrangement within the polypeptide chain. This is called the secondary structure and exists in two main forms.
- (i) State the two forms of secondary structure found in proteins.
- (ii) Draw a diagram to illustrate **one** form of secondary structure.

(c) There are around 20 different common amino acids found in humans most of which have the same general structure.

The nature of the group R affects which bonds are formed as the secondary structure of the protein is further folded to give the tertiary structure.

Complete the table indicating the type of **tertiary** bonding that each pair of the amino acid residues is likely to produce.

residue 1	residue 2	type of tertiary bonding
-HNCH(CH ₂ CH ₂ CH ₂ CH ₂ NH ₂)CO-	-HNCH(CH ₂ CH ₂ CO ₂ H)CO-	
-HNCH(CH ₃)CO-	-HNCH(CH ₃)CO-	
-HNCH(CH ₂ SH)CO-	-HNCH(CH ₂ SH)CO-	
-HNCH(CH ₂ OH)CO-	-HNCH(CH ₂ CO ₂ H)CO-	

(Nov 2011 P41 Q6)

- Q4 The formation of proteins is a key process in the growth and repair of tissues in living organisms.
- (a) (i) Study the structures of the three molecules below. One of the molecules could be a building block for a protein while the other two could be building blocks for other biological polymers.

Which of the three could be a building block for a protein? Explain your answer.

(ii) For which biological polymer could **one** of the other molecules form a building block?

molecule polymer

- **(b)** Protein molecules have four levels of structure as the long molecules fold and take shape.
- (i) The primary structure is the sequence of amino acids in the protein chain. What type of bonding exists between the amino acids in this chain?

(ii) What type of bonding can exist in all of the other types of structure?
(iii) Name one type of bonding that does not occur in the primary or secondary structure of the protein.
(c) Many proteins play an important role in catalysing chemical reactions in living organisms. (i) What name is given to these catalysts?
(ii) Give two changes in conditions under which these catalysts may be inactivated, explaining the chemical reason for this in each case.
(Nov 2011 P43 Q6)
Q5 In key reactions responsible for growth and repair in the human body, amino acids react together to form polymers known as proteins. (a) (i) What <i>type of reaction</i> is this polymerisation?
(ii) From stocks of glycine and alanine, it is possible to make the dipeptide gly-ala. Using the same three-letter abbreviations for the amino acids, give the structures of all other possible dipeptides that can be made from these stocks of amino acids. [3]
(b) (i) DNA consists of a double helix formed by two strands held together by hydrogen bonds between base pairs. Sketch a section of DNA showing two base pairs, using blocks for the various components. You should label all of the components.
(ii) Suggest what the effect on DNA replication would be if the hydrogen bonds between the strands were replaced by stronger bonds, e.g. covalent bonds.
[4]

change in the	e triplet code hy some cha	as sickle-cell anaemia, are c nges in the triplet code do no	·	
		in the tertiary structure of a position of the distribution of the	protein would result from a	mutation
		O H ₂ N—CH—C—OH I CH ₂ I C=O OH	I—CH ₂ —CO ₂ H	
		aspartic acid	glycine	
Q6 (a) The ta		ne structures of four amino action by indicating the type of terti	cids found in proteins in the	
likely to have		ino acid is present in a prote	T	П
	amino acid	structure	type of interaction	+
	alanine	H ₂ NCH(CH ₃)CO ₂ H		+
	cysteine	H ₂ NCH(CH ₂ SH)CO ₂ H		+
	lysine serine	H ₂ NCH((CH ₂) ₄ NH ₂)CO ₂ H		+
following me there.	s play an imp tal ions, outli	H ₂ NCH(CH ₂ OH)CO ₂ H portant role in the biochemist ne one of the places in the bo	ody it can be found and its	main role
potassium				

zinc					
energy is	chemical reactions at a cellular level rec s largely provided by the breakdown of or an equation showing the breakdown of th	ne particular compound.	his		
(ii) What	type of chemical reaction is this?				
in the procell mem People w • water is outside to • their sy Based or (i) sugge	(d) Cystic fibrosis is a genetic disease caused by a mutation in the DNA sequence resulting in the production of a faulty version of an important protein which acts as an ion pump in the cell membrane. This pump controls the flow of ions into and out of cells. People with the faulty protein show two major symptoms. • water is retained in cells in the lungs resulting in the formation of a thick, sticky mucous outside the cells; • their sweat is very salty. Based on the information given for people with cystic fibrosis, (i) suggest which ions are involved in the ion flow, (ii) suggest and explain what type of bonding might result in thick or sticky mucous.				
		(June 2012 F			
three-din (a) Study	eins are complex molecules made up from nensional structure. If the table which describes aspects of bo type, indicate whether it contributes to the	nding in proteins. For each descripti	ion of a		
	bonding type	structure involved			
	disulfide bonds between parts of the chain				
	hydrogen bonds in a β-pleated sheet				
	ionic bonds between parts of the chain				
	peptide links between amino acids				

BIOCHEMISTRY

(b) Explain, with the use of diagrams as appropriate, the difference between competitive and non-competitive inhibition of enzymes.

Application

Application	BIOCHEMIS	TRY
		[4]
labels, the bonds holding th	strand of DNA. Draw a matchine two strands together. Name e bonds to each base in the stranger phosphate	
	T	G
names of bases		
		(Nov 2012 P42 Q6)
Q8 The proteins in the hum amino acids. Alanine is a ty		s made up of around 20 different
	н Н о	
	N—C—C	
	H CH OH	
	OI 13	
(a) Glycine H2NCH2CO2H	alanine is the simplest amino acid an	d differs from each of the other
	nt way. What is this difference?	
		[1]
shapes. This is referred to a		with complex three-dimensional tructures of a protein.
structure		
(ii) Give two examples of be responsible in each case.	onding causing the tertiary str	ructure, and give the amino acid
bonding	amino acid	
(c) Suggest why globular pr	oteins, such as enzymes, cont ompared to the amounts of sor	[6] tain relatively small amounts of me other amino acids. You may

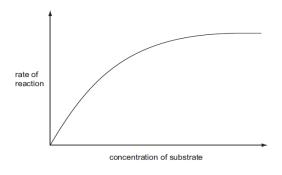
Application				E	ВІОСНЕ	MIST	RY				
		•••••		•••••		•••••		•••••	•••••	•••••	
(d) DNA consists of a with one of four bases to the sugar. (i) The two strands of	s – ad	enine	(A), cyt	osine	(C), gu	anine	(G) and	thym	nine (T)	– attac	
pairs of bases. What a	are the	e pairs	s of bas	es?							
In protein synthesis, s the ribosome in order three bases codes for The codes are summa	to ass	sembl amino	e the ar acid, w	nino a	acids for	the n	ew prot	tein cl	nain. E	ach gro	
	UUU UUC UUA UUG	phe phe leu leu	UCU UCC UCA UCG	ser ser ser	UAU UAC UAA UAG	tyr tyr stop stop	UGU UGC UGA UGG	cys cys stop trp			
	CUU CUC CUA CUG	leu leu leu leu	CCU CCC CCA CCG	pro pro pro pro	CAU CAC CAA CAG	his his gln gln	CGU CGC CGA CGG	arg arg arg arg			
	AUU AUC AUA AUG	ile ile ile met/ start	ACU ACC ACA ACG	thr thr thr thr	AAU AAC AAA AAG	asn asn lys lys	AGU AGC AGA AGG	ser ser arg arg			
	GUU GUC GUA GUG	val val val val	GCU GCC GCA GCG	ala ala ala ala	GAU GAC GAA GAG	asp asp glu glu	GGU GGC GGA GGG	gly gly gly			
(ii) The coding for all p codes shown in the ta produce?		/hat a	imino ad	cid se	quence	would	the fol				•
		-AUC	GGGUA	GCC	UCGCA	UCG	JAA-				
(iii) What would be the base at position 10 in								tation	that ch	anged	the
									(Nov 2	 2012 P4	3 Q6)
Q9 There are two imp synthesis and the forn (a) Complete the table	nation	of DI	NA.						nisms -	- proteir	·

each substance could be used.

substance	protein synthesis	formation of DNA
adenine		
alanine		
aspartate		
phosphate		

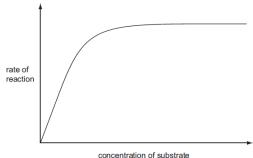
• •		ent helical structures. nese helical structures	Briefly describe the bos.	onding that
protein				
DNA				
	uld include refer		and <i>tertiary</i> structures re of the bonding and	
			(Jur	ne 2013 P41 Q6)
synthesis and the	e formation of DI table by placing	NA.	ur within living organis column to indicate in w	·
,	substance	protein synthesis	formation of DNA	
	cysteine			
,	cytosine			
,	glutamine			
	guanine			
(b) DNA consists (i) Describe the b strand is joined b	onding betweer		NA and state which pa	art of each
/ii) How doos the	atropath of this	handing ralets to the	machaniam of the ron	lication of DNA
(ii) now does the	sarengin or inis	bonding relate to the	mechanism of the rep	ilication of DINA!
(c) Some disease basis of these ch			ture of proteins. Expla	

Application		BIOCHEMISTR	Y
			(June 2013 P42 Q6)
044 (a) Duatain va	-l		,
	olecules are forme amino acids are g		of amino acids in the body. The
	_	O	0
шь	ОН	но	ОН
H ₂ N	'	NH ₂	NH ₂
	glycine (gly)	serine (ser)	valine (val)
(i) How many different acids shown?	rent tripeptides car	n be made using one mo	plecule of each of the amino
acius silowii:			
(ii) Draw the triper	ntide ser-alv-val sh	owing the peptide bonds	s in displayed form
(ii) Diaii ale alper	ondo don giy van, di	iowing the populae bende	o in alopiayou form.
(iii) Within the trip	eptide, which amin	o acid provides a hydrop	phobic side chain?
	•	, , , , ,	
		ds giving proteins their s	
		, state two types of bond ved in this bonding.	ding that can be formed and the
		·	
bond		groups	
			emical reactions. The efficiency les known as inhibitors. Explain
how both competis		etitive inhibitors prevent	
efficiently. (i) competitive inhibits	ibitors		
(ii) non-competitiv	e inhibitors		
(iii) The graph sho	ows the rate of an e	enzyme-catalysed reaction	on against the substrate
concentration in th	ne absence of an in	hibitor. On the same axe	es, sketch a graph showing the
rate of this reactio	n it a <i>non-competit</i>	ive inhibitor was present	



(Nov 2013 P42 Q6)

- Q12 (a) Enzymes are particular types of proteins that catalyse chemical reactions. The efficiency of enzymes can be reduced by the presence of other substances known as inhibitors.
- (i) State **one** example of a substance that can act as a *non-competitive* inhibitor in enzyme reactions.
- (ii) For the inhibitor you have identified, explain why it is a non-competitive inhibitor.
- (iii) The graph shows the rate of an enzyme-catalysed reaction against the substrate concentration in the absence of an inhibitor. On the same axes, sketch a graph showing the rate of this reaction if a *competitive inhibitor* was present.



- **(b)** DNA is responsible for encoding the amino acid sequence to produce proteins. Ribosome, tRNA and mRNA are all involved in the process of protein synthesis.
- (i) Write ribosome, tRNA and mRNA in the boxes below to show the correct sequence in which they are involved.



- (ii) Sequences of three bases code for specific amino acids. The code UGA however does not usually code for an amino acid. Suggest its use.
- (c) Much of the energy used in biochemical reactions is provided by the hydrolysis of the molecule ATP
- (i) What are the breakdown products of the hydrolysis of ATP?
- (ii) Give two uses for the energy released by ATP hydrolysis in cells.

1.