

Question			Answer	Mark	Guidance
1	(a)	(	polar <b>and</b> brown bear ;	1	
		(ii)	<i>no because</i> one, more closely related to / in same group as , raccoons <b>and</b> one , to / with, bears / AW ;	1 max	<b>DO NOT CREDIT</b> answer if in context of yes
	(b)	(	knowledge , tentative / uncertain / subject to change ; to re-test / check, hypotheses / results ;	2	<b>IGNORE</b> incomplete, new technology <b>IGNORE</b> to validate
		(ii)	<p><b>1</b> <i>idea that</i> haemoglobin could be , an <u>adaptation</u> (to the environment) / an <u>adaptive</u> feature ;</p> <p><b>2</b> <i>idea that</i> low oxygen partial pressure is selective agent <b>or</b> both subject to the same selection pressure ;</p> <p><b>3</b> (haemoglobin of both) has high oxygen affinity / dissociation curve shifted to left ;</p> <p><b>4</b> convergence / similarity not due to shared ancestry ;</p>	3 max	<p><b>3</b> <b>ACCEPT</b> haemoglobin can uptake O<sub>2</sub> at low partial pressure</p> <p><b>4</b> <b>ACCEPT</b> description e.g. "changes happen to both independently" <b>IGNORE</b> "red and giant panda may not be closely related" (as given in question)</p>

Question		Answer	Mark	Guidance
	(c)	<p><b>step 2</b> PCR / polymerase chain reaction ;</p> <p><b>step 3</b> genetic modification / genetic engineering ;</p> <p><b>step 4</b> electrophoresis ;</p>	3	<p><b>FA on each line</b></p> <p><b>ACCEPT</b> gene cloning / transformation</p> <p><b>ACCEPT</b> (gel) chromatography</p>
	(d)	<p>triplet code <b>or</b> 3 bases = 1 amino acid ;</p> <p>525 ;</p> <p>3 bases are , stop / (chain) termination , codon ;</p>	3	<b>DO NOT CREDIT</b> triplet makes amino acid
	(e)	( ox ;	1	<b>FA</b>
	(ii)	<p><b>1</b> genetic code is degenerate ;</p> <p><b>2</b> more than 1, triplet / codon, for same amino acid ;</p> <p><b>3</b> silent / neutral, mutations ;</p> <p><b>4</b> <i>idea that</i> DNA, changes more than / is more different to, protein ;</p>	3 max	<p><b>1 ACCEPT</b> redundant</p> <p><b>2 DO NOT CREDIT</b> 'make' the same amino acid</p> <p><b>4 ACCEPT</b> polypeptide / amino acid sequence <b>ACCEPT</b> nucleotide sequence for DNA</p>
		<b>Total</b>	<b>17</b>	

Question			Expected Answer	Mark	Additional Guidance
2	(a)	(i)	microbes / (living) organisms / cells / enzymes ;  (make) product / for human benefit / (carry out) conversion / reaction / industrial process ;	2	<b>CREDIT</b> microorganisms / bacteria / prokaryotes / fungi <b>CREDIT</b> living things <b>CREDIT</b> cell components / parts of cells  <b>CREDIT</b> example such as (named) food or medicine BUT <b>IGNORE</b> cheese (as stated in question) <b>IGNORE</b> process unqualified
2	(a)	(ii)	microbes / <b>AW</b> , killed / removed / not present ;  enzymes <u>denatured</u> ;  (so no) competitors / unwanted reactions / (human) health risk ;	2 max	<b>Mark the first two suggestions</b> <b>IGNORE</b> contamination / sterile <b>IGNORE</b> idea of preserving milk  <b>AW</b> for microbes as in (a)(i) plus <b>ACCEPT</b> organisms  <b>DO NOT CREDIT</b> microbes denatured  <b>CREDIT</b> (no) competition <b>CREDIT</b> (no) food spoilage / change of flavour / loss of quality <b>CREDIT</b> (no) pathogens / harmful microbes / TB  <b>“Kills harmful microbes”</b> or <b>“Kills pathogens”</b> scores 2 marks (mps 1 & 3)

Question			Expected Answer	Mark	Additional Guidance
2	(b)	(i)	<p>1 enzyme ;</p> <p><i>plus any 2 of the following</i></p> <p>2 (enzyme) not, changed / used up ; <b>ora</b></p> <p>3 <i>idea of ESC (forms) / substrate and enzyme (bind) ;</i></p> <p>4 products (and enzyme) released at end ;</p>	<p>1</p> <p>max 2</p>	<p><b>Award mp 1 plus 2 max from the other mark points</b></p> <p>1 <b>ACCEPT</b> globular / tertiary / catalyst / catalytic (protein)</p> <p>2 <b>ora</b> = can be used again / re-used <b>IGNORE</b> enzyme recycled</p> <p>3 ESC = enzyme-substrate complex <b>ACCEPT</b> substrate entering active site</p>
2	(b)	(ii)	<p>1 (enzyme can be removed to be) used again ;</p> <p>2 (enzyme can) to leave pure(r) product ; <b>ora</b></p> <p>3 (enzyme) more stable / more efficient / works better ;</p>	<p>2</p>	<p><b>Mark the FIRST suggestion on each numbered line</b> <b>IGNORE</b> 'cheaper' without qualification</p> <p>2 <b>ACCEPT</b> cheaper / easier, downstream processing</p> <p>3 <b>CREDIT</b> less susceptible to, pH / temperature, change / extremes  <b>"enzymes work at high temperatures"</b> = 0  <b>"enzymes work at higher temperatures"</b> = 1  (because comparative statement made)</p>

Question		Expected Answer	Mark	Additional Guidance
2	(c)	<p><b>This is a QWC question</b></p> <p><i>Section I - Obtaining the gene</i></p> <p>1 use restriction, enzyme / endonuclease ;</p> <p>2 to, cut out / get / isolate, (rennin) gene / DNA coding for rennin or to, fragment / digest, DNA ;</p> <p>3 <u>gene</u> probe ; <b>OR</b></p> <p>4 obtain rennin mRNA ;</p> <p>5 (use) reverse transcriptase ;</p> <p>6 to make cDNA ; <b>OR</b></p> <p>7 sequence, rennin (protein) ;</p> <p>8 work out base code ;</p> <p>9 make this DNA sequence ;</p> <p>10 sticky ends ;</p> <p><i>Section II - Vector</i></p> <p>11 cut (open), plasmid / phage ;</p> <p>12 using same <u>restriction</u> enzyme ;</p> <p>13 annealing / base pairing of sticky ends ;</p> <p>14 join sugar-phosphate backbones ;</p> <p>15 (using DNA) ligase ;</p> <p>16 <u>recombinant</u>, vector / plasmid / phage / DNA ;</p> <p><i>Section III - Introduction into host cell</i></p> <p>17 mix with bacteria ;</p> <p>18 detail of conditions ;</p> <p>19 <u>transformation</u> (plasmid) / <u>transduction</u> (phage) ;</p>	max 7	<p>1 <b>CREDIT</b> named example e.g. <i>Eco R1, Bam H1, Hin dIII</i></p> <p>2 <b>DO NOT CREDIT</b> 'cut gene' <b>IGNORE</b> 'break up DNA'</p> <p><b>NOTE</b></p> <p>1-9 <b>CREDIT</b> whichever of the three alternative "obtaining the gene" protocols yields most marks, <b>either award marking points</b></p> <p>1- or 4-6 or 7-9</p> <p>10 can be awarded, once only, in Sections I or II</p> <p>11 <b>DO NOT CREDIT</b> 'cut out plasmid' <b>DO NOT CREDIT</b> 'ring of DNA' unless it is clear that plasmid is being referred to</p> <p>12 <b>CREDIT</b> same named enzyme (re. mp1)</p> <p>13 <b>CREDIT</b> idea of sticky end bases hydrogen bonding</p> <p>14 <b>CREDIT</b> formation of phosphodiester bonds</p> <p>18 e.g. Ca<sup>2+</sup> ions added / heatshock (freeze then inc to 40°C)</p> <p>19 <b>CREDIT</b> transform / transformed / transduce / transduced <b>IGNORE</b> transgenic</p>
		<b>QWC – sequencing of steps – at least 1 mark point scored from each of the three sections, in the correct order ;</b>	1	<p>I. obtaining gene (mp 1 – 9) followed by</p> <p>II. vector (mp 13 – 16) followed by</p> <p>III. introduction to host cell (mp 17 – 19)</p>
<b>TOTAL</b>			<b>17</b>	

Question		Expected Answers		Marks	Additional Guidance									
3	(a)		<table border="1"> <thead> <tr> <th></th> <th>similarity</th> <th>difference</th> </tr> </thead> <tbody> <tr> <th>structure</th> <td>mitochondria <b>or</b> vesicles <b>or</b> postsynaptic receptors ;</td> <td>NMJ membrane(s), wavy / AW * <b>ora</b> <b>or</b> receptors different (shape) <b>or</b> enzymes in different places ;</td> </tr> <tr> <th>function</th> <td>(neuro)transmitter, released / crosses gap <b>or</b> changes potential difference / AW ** <b>or</b> enzymes break down (neuro)transmitter ;</td> <td>different neurotransmitters / ACh vs. dopamine <b>or</b> muscle contraction vs. nerve impulse <b>or</b> different enzymes ;</td> </tr> </tbody> </table>		similarity	difference	structure	mitochondria <b>or</b> vesicles <b>or</b> postsynaptic receptors ;	NMJ membrane(s), wavy / AW * <b>ora</b> <b>or</b> receptors different (shape) <b>or</b> enzymes in different places ;	function	(neuro)transmitter, released / crosses gap <b>or</b> changes potential difference / AW ** <b>or</b> enzymes break down (neuro)transmitter ;	different neurotransmitters / ACh vs. dopamine <b>or</b> muscle contraction vs. nerve impulse <b>or</b> different enzymes ;		<p><b>One mark per box</b></p> <p><i>difference</i> <b>NMJ</b> is neuromuscular junction * <b>AW A CEPT</b> wiggly / bumpy / not smooth / rough / larger SA / any suitable description <b>but IGNORE</b> microvilli</p> <p><i>difference</i> <b>ACh</b> is acetylcholine</p> <p><i>similarity</i> ** <b>AW CREDIT</b> depolarises / -70 mV → +40 mV <b>but IGNORE</b> pass on action potential</p>
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3	(b)	(i)	<p>1 phenelzine ;</p> <p><b>no ecf from incorrect drug</b></p> <p>2 <i>idea that</i> does not bind to (dopamine) receptor ; <b>ora</b></p> <p>3 <i>idea that</i> binds to, MAO / enzyme ;</p> <p>4 allosteric site / non-competitive inhibitor ;</p>	1	<p><b>Award mp1 and, if correct, any 1 from the remaining points</b></p> <p>2 <b>CREDIT</b> other two do bind to dopamine receptor</p> <p>3 <b>IGNORE</b> inhibits, MAO / enzyme (as given in the question)</p> <p>4 <b>ACCEPT</b> “not a competitive inhibitor”</p>									
				max 1										
3	(b)	(ii)	(drug) occupies / blocks / binds to, (dopamine) receptors ; without causing, action potential / response ; reduces <b>effect of</b> dopamine / is a dopamine antagonist ;	2	<p><b>CREDIT</b> “without causing depolarisation” / AW <b>DO NOT CREDIT</b> “inhibits dopamine” or “reduces dopamine levels”</p>									

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3	(c)	(i)	humans are, diploid / $2n$ ; chromosomes, are in pairs / homologous ; one, (copy / gene / allele), from each parent / on each chromosome of pair ;	2 max	<b>DO NOT CREDIT</b> ref to bivalents
3	(c)	(ii)	(gel) <u>electrophoresis</u> ;	1	
3	(d)	1	13 b-p deletion (has most serious consequences) ;	3 max	6 <b>CREDIT</b> could be a silent mutation / 1 b-p substitution may not have an effect
		2	frameshift / alter reading frame ;		
		3	genetic code is triplet / read in groups of 3 bases ;		
		4	alters all amino acids (coded for) after the mutation ;		
		5	21 b-p deletion causes 7 amino acids to be lost ;		
		6	substitution changes, one / no, amino acids ;		
3	(e)	1	<u>natural selection</u> ;	4 max	3 <b>CREDIT</b> increases reproductive success / AW 4 <b>ACCEPT</b> more promiscuous / AW  6 <b>MUST HAVE</b> time element
		2	<u>selective advantage</u> ;		
		3	(allele / behaviour) increases, survival / breeding / AW ;		
		4	(because) helped, find food / find new resources / make new tools / get mates ;		
		5	<u>allele</u> passed on (to next generation) ;		
		6	(allele / behaviour) increased in frequency over, generations / time ;		
<b>Total</b>				<b>18</b>	