

| Question | | | Answer | Mark | Guidance |
|----------|-----|-----|-------------------------|------|--|
| 1 | (a) | | (works) outside cells ; | 1 | ACCEPT secreted / AW , from cells ACCEPT works in named extracellular environment e.g. digestive tract IGNORE doesn't work in cells |
| | (b) | (i) | time / time taken ; | 1 | Mark the first answer. If the answer is correct and another answer is given that is incorrect or contradicts the original answer, then = 0 marks ACCEPT 'how long it took ...' |

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| (b) | (ii) | <p><i>linear part of the graph means...</i></p> <p>1 <u>more</u> (successful) collisions with (amylase) <u>active site</u> (at increasing starch concentration) ; ora</p> <p>2 <u>more</u> ESC (at increasing starch concentration) ; ora</p> <p>3 so more product formation in a <u>given time</u> (at increasing starch concentration) ; ora</p> <p><i>curve / plateau , means...</i></p> <p>4 all / most , <u>active sites</u> (of amylase) are occupied ;</p> <p>5 enzyme / amylase , working , at / near, maximum rate / V_{max} ;</p> <p>6 (so) further increase in starch concentration has no effect (on rate) ;</p> <p>7 enzyme <u>concentration</u> , is / becomes , <u>limiting</u> factor ;</p> | 5 max | <p>ACCEPT glucose / maltose for product throughout ACCEPT substrate for starch throughout</p> <p>1 ACCEPT few(er) active sites occupied at low starch concentrations</p> <p>2 ACCEPT ESC formed more easily</p> <p>3 AWARD only if linked to the context of marking points 1 or 2 e.g. 'more product formation in a given time because of more collisions with the enzyme' gets mp3 but not mp1 because active site not mentioned 3 IGNORE <u>rate</u> as this is a description of graph</p> <p>4 ACCEPT all active sites are full of substrate</p> <p>5 ACCEPT enzyme at full capacity</p> <p>6 Must link to 4 or 5 6 AWARD only if mp 4 or 5 given 6 DO NOT CREDIT rate decreases</p> <p>7 ACCEPT the increasing part of the graph is because starch <u>concentration</u> is the <u>limiting</u> factor</p> |

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| (b) | (iii) | <p>1 (so) charges in active site do not change ; ora</p> <p>2 (so) hydrogen / ionic , bonds unaffected ; ora</p> <p>3 (so) tertiary structure / 3D shape / active site , unaltered ; ora</p> <p>4 (so) enzyme / tertiary structure , does not <u>denature</u> ; ora</p> <p>5 (so) substrate , fits / is complementary shape to , <u>active site</u> ; ora</p> <p>6 so the results are <u>valid</u> / as the <u>rate</u> (of reaction) will vary if pH varies / so that only one (independent) variable is changed ;</p> | 3 max | <p>The mark points refer to a constant pH preventing damage to the enzyme. CREDIT throughout the appropriate marking point for an answer that describes what would happen if the pH changed.</p> <p>2 DO NOT CREDIT peptide / disulphide , bonds break</p> <p>2 DO NOT CREDIT in context of heat / vibration</p> <p>2 IGNORE hydrophobic / hydrophilic</p> <p>3 IGNORE ref to denaturing active site</p> <p>3 IGNORE tertiary structure breaks</p> <p>3 ACCEPT tertiary structure affected</p> <p>3 Cannot be inferred from mp5 – must be stated</p> <p>4 IGNORE ref to denaturing active site</p> <p>4 DO NOT CREDIT kill / die</p> <p>5 IGNORE enters / binds with</p> <p>6 IGNORE fair test / reliable / accurate</p> |

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|--|---|--|---------|--|---------------|-------------------|---|--|--|---|---|---|------------------------|--------------------------|--|--|---|--|
| (b) | (iv) | <p>temperature (of the reaction mixture) ; enzyme / amylase , concentration ;</p> <p>(total) volume of (reaction) <u>solution</u> ;</p> <p>concentration of , cofactors / chloride ions / Cl⁻ ;</p> | 2 max | <p>Mark the first answer on each prompt line. If the answer is correct and another answer is given that is incorrect or contradicts the original answer, then = 0 marks DO NOT CREDIT substrate / starch , concentration (as this is the independent variable)</p> <p>DO NOT CREDIT amount</p> <p>ACCEPT volume of enzyme solution DO NOT CREDIT amount</p> <p>ACCEPT concentration of coenzymes</p> <p>IGNORE time / agitation / inhibitors</p> | | | | | | | | | | | | | | |
| (c) | (i) | <table border="1" style="width: 100%; border-collapse: collapse;"> <thead> <tr> <th style="width: 50%; text-align: center;">Amylose</th> <th style="width: 50%; text-align: center;">Cellulose</th> </tr> </thead> <tbody> <tr> <td style="text-align: center;"><i>coiled</i></td> <td style="text-align: center;"><i>no coiling</i></td> </tr> <tr> <td style="text-align: center;">(contains) α / alpha / A / a , -glucose</td> <td style="text-align: center;">(contains) β / beta / B / b , -glucose ;</td> </tr> <tr> <td style="text-align: center;">α / alpha / A / a 1-4 glycosidic bonds</td> <td style="text-align: center;">β / beta / B / b 1-4 glycosidic bonds ;</td> </tr> <tr> <td style="text-align: center;">all , monomers / AW , in same orientation</td> <td style="text-align: center;">alternate monomers at , 180° / AW , to each other ;</td> </tr> <tr> <td style="text-align: center;">granular / not fibrous</td> <td style="text-align: center;">fibrous / not granular ;</td> </tr> <tr> <td style="text-align: center;">H bonds within molecule / no (H) bonds (between molecules)</td> <td style="text-align: center;">(H) bonds between adjacent molecules ;</td> </tr> </tbody> </table> | Amylose | Cellulose | <i>coiled</i> | <i>no coiling</i> | (contains) α / alpha / A / a , -glucose | (contains) β / beta / B / b , -glucose ; | α / alpha / A / a 1-4 glycosidic bonds | β / beta / B / b 1-4 glycosidic bonds ; | all , monomers / AW , in same orientation | alternate monomers at , 180° / AW , to each other ; | granular / not fibrous | fibrous / not granular ; | H bonds within molecule / no (H) bonds (between molecules) | (H) bonds between adjacent molecules ; | 3 | <p>Mark the first 3 responses AWARD 1 mark for each correct row irrespective of boxes Three correct rows of responses written within the same box can be awarded 3 points.</p> <p>ACCEPT every second one is flipped</p> <p>ACCEPT fibres / microfibrils / fibrils / macrofibrils DO NOT CREDIT myofibrils ACCEPT grains</p> <p>ACCEPT '(cross)links' as AW for 'bonds'</p> |
| Amylose | Cellulose | | | | | | | | | | | | | | | | | |
| <i>coiled</i> | <i>no coiling</i> | | | | | | | | | | | | | | | | | |
| (contains) α / alpha / A / a , -glucose | (contains) β / beta / B / b , -glucose ; | | | | | | | | | | | | | | | | | |
| α / alpha / A / a 1-4 glycosidic bonds | β / beta / B / b 1-4 glycosidic bonds ; | | | | | | | | | | | | | | | | | |
| all , monomers / AW , in same orientation | alternate monomers at , 180° / AW , to each other ; | | | | | | | | | | | | | | | | | |
| granular / not fibrous | fibrous / not granular ; | | | | | | | | | | | | | | | | | |
| H bonds within molecule / no (H) bonds (between molecules) | (H) bonds between adjacent molecules ; | | | | | | | | | | | | | | | | | |

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| | (c) (ii) | (tensile) strength / strong ; (H) bonds / links , can form (between adjacent fibrils) ; insoluble ; | 2 max | ACCEPT mechanical strength IGNORE fibrous / rigid ACCEPT fibres / microfibrils / fibrils / macrofibrils IGNORE refs to bonding with water IGNORE ionic / myofibrils ACCEPT crosslinks DO NOT CREDIT peptide / covalent / glycosidic / disulfide etc |
| | | Total | 17 | |

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|----------|-----|------|--|-------|--|
| 2 | (a) | (i) | udder size / milk production / meat production / growth rate / muscle (as proportion of body mass) ; | 1 | ACCEPT number of offspring per birth IGNORE unqualified references to size IGNORE references to , horns / placidity , unless the answer links this with more energy diverted to productivity |
| | (a) | (ii) | <p>1 artificial <u>selection</u> ;</p> <p>2 (selection of) named desired feature (linked to productivity) ;</p> <p>3 (cross)breed , selected / AW , cattle ;</p> <p>4 (cross)breed, best / selected / AW, offspring ;</p> <p>5 over (many) generations ;</p> | 4 max | <p>1 IGNORE 'selective breeding' as mentioned in part (i)</p> <p>2 ACCEPT e.g. weigh them / measure them / see who produces the most milk / choose the biggest / udder size 2 IGNORE select the best 2 CREDIT marker assisted selection / progeny testing 2 DO NOT CREDIT if clearly not in the context of selective breeding, e.g. change their diet to make them produce more milk'</p> <p>3 ACCEPT 'parents' as AW for 'cattle' 3 ACCEPT 'reproduce / mate / <u>inter</u>breed' as AW for 'breed' 3 DO NOT CREDIT inbred</p> <p>2&3 'breed cattle with high milk productivity = 2 marks</p> <p>4 IGNORE 'crossbreed offspring' without qualification. Answer must imply some selection of offspring.</p> <p>5 DO NOT CREDIT few 5 ACCEPT several</p> |
| | (b) | (i) | (contains) all / each , of , nutrients / food groups ; in correct proportions / AW ; | 2 | ACCEPT a list of food groups that contains at least – protein, fat, carbohydrate, vitamins, minerals IGNORE components ACCEPT right amount of |

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|----------|----------|---|------|---|
| | (b) (ii) | <p>A glycerol ;</p> <p>C <u>unsaturated</u> fatty acid ;</p> <p>D <u>ester</u> , bond / link ;</p> | 3 | <p>Mark the first answer on each prompt line. If the answer is correct and another answer is given that is incorrect or contradicts the original answer, then = 0 marks</p> <p>A IGNORE molecule</p> <p>C ACCEPT unsaturated hydrocarbon , tail / chain</p> <p>D IGNORE covalent</p> |

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| | (b) (iii) | <p>1 contains , large amounts of energy / more energy than individual needs ;</p> <p>2 increased , fat / lipid , deposition / storage ;</p> <p>3 (associated with) <u>obesity</u> ;</p> <p>4 (lots of meat and dairy in diet could mean) lack of <u>other (named) food groups</u> / AW ;</p> | 3 max | <p>1 ACCEPT contains , too many calories / excess energy</p> <p>1 ACCEPT contains a lot of <u>saturated</u> fat</p> <p>2 ACCEPT in context of arteries and adipose tissue</p> <p>2 ACCEPT cholesterol / LDL as AW for fat</p> <p>2 IGNORE build up</p> <p>3 IGNORE CHD (as not malnutrition)</p> <p>4 ACCEPT nutrients as AW for food groups</p> <p>4 IGNORE unbalanced diet</p> <p>4 IGNORE fat / protein</p> |
| | (c) | <p>1 reduces , water potential / Ψ , outside , microbial / bacterial / fungal , cells ;</p> <p>2 (microbes) lose water and cannot , reproduce / survive / carry out metabolic reactions / AW ;</p> <p>3 water moves by osmosis ;</p> | 3 | <p>1 Cannot be implied from references to water potential gradient</p> <p>1 ACCEPT reduces beef water potential</p> <p>1 IGNORE solute potential</p> <p>1 IGNORE viruses</p> <p>2 ACCEPT bacteria lose water and die</p> <p>2 AWARD only in context of microbes dehydrating</p> <p>2 IGNORE viruses</p> <p>2 IGNORE beef losing water so microbes cant reproduce</p> <p>3 ACCEPT in any correct water potential context</p> |
| Total | | | 16 | |

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|----------|-----|-------|--|-------|---|
| 3 | (a) | (i) | blue-black / black / dark blue ; | 1 | ACCEPT dark purple / purplish-blue DO NOT CREDIT blue or purple unqualified by darkness ACCEPT acceptable colour change |
| | (a) | (ii) | 1 between oxygen and hydrogen (atoms) ; 2 (between) electronegative / δ^- , and electropositive / δ^+ ; | 2 | CREDIT marking points from clearly labelled diagram max 1 if incorrect charges are on atoms 1 DO NOT CREDIT molecules / ions 2 DO NOT CREDIT ions / + and – 2 ACCEPT slight / partial (negative / positive), charge |
| | (a) | (iii) | 1 hydrogen / H, bonds break ; 2 <u>helix</u> , lost / unravels / AW ; 3 iodine, released / no longer in complex / AW ; | 2 max | IGNORE refs to denaturation 2 ACCEPT spiral / coil 3 ACCEPT no longer contained in helix |

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|----------|--|---------------------|--|
| (b) | <p>1 take samples at a range of times / AW ;</p> <p>B2 same <u>volumes</u> (of solutions) added / removed (each time) ;</p> <p>B3 heat with, Benedict's (solution) / CuSO₄ and NaOH ;</p> <p>B4 (use of) excess Benedict's ;</p> <p>B5 changes to, green / yellow / orange / brown / (brick) red ;</p> <p>C6 remove precipitate / obtain filtrate ;</p> <p>C7 colorimeter ;</p> <p>8 calibrate / zero, using, a blank / water / (unreacted) Benedict's ;</p> <p>9 use (red / orange) filter ;</p> <p>T10 reading of, transmission / absorbance OR mass of precipitate ;</p> <p>11 more transmission / less absorbance, of filtrate, OR greater mass ppt, = more maltose present ; ora</p> <p>12 using, standard / known, concentrations (of maltose) ;</p> <p>13 (obtain) <u>calibration</u> curve ;</p> <p>14 <u>plot</u>, transmission / absorbance / mass of ppt, against (reducing sugar) concentration ;</p> <p>15 <u>use graph</u> to read off concentration of maltose / AW ;</p> | <p>6 max</p> | <p>B2 must be in context of Benedict's test rather than reaction mixture</p> <p>B3 DO NOT CREDIT boil / warm</p> <p>B3 DO NOT CREDIT if Benedict's added to the mixture at the beginning</p> <p>C6 CREDIT description of method e.g. filtering / centrifuging / decanting</p> <p>8 IGNORE 'control'</p> <p>9 DO NOT CREDIT if colour of filter is incorrect</p> <p>T10 ACCEPT 'measure how much light, does / does not, pass through'</p> <p>11 if unfiltered Benedict's / precipitate is clearly indicated as being present in sample, ACCEPT 'less transmission / more absorbance, = more maltose present'</p> <p>11 DO NOT CREDIT if precipitate is added to colorimeter</p> <p>12 CREDIT 'serial dilutions'</p> |
| | QWC – correct sequence ; | 1 | 1 of mps B2 to B5 , then mp C6 or C7 , then mp T10 |

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|---|---|---|---------|--|---|---------|---------|---------|---------|--|--|--|--|--|---------|---------|---------|---------|---------|---------|---------|---------|---------|-----------------------|---|----|----|----|----|----|----|-----|-----|----------------------|---|----|----|----|----|----|----|----|----|---|---|----|----|----|----|----|----|
| | (c) (i) | <p>1 increases / greater / faster ;</p> <p>2 reaction completed in / plateaus after / concentration is 100% after, 3.5 minutes ;</p> <p>3 figures with units to support mp 1 ;</p> | 2 max | <p>1 ACCEPT any time between 3.45 and 3.55 min.</p> <p>3 two maltose concentrations (+ or – chloride) for a given time or two times (+ or – chloride) for given maltose concentration.</p> <p>3 ACCEPT calculated difference</p> <p>3 DO NOT CREDIT if ‘%’ and ‘min.’ not given</p> <p>3 ACCEPT any concentration within $\pm 1\%$ and time within ± 0.05 min.</p> | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | | <table border="1"> <thead> <tr> <th rowspan="2">Presence or absence of chloride ions</th> <th colspan="10">The percentage concentration of maltose (%) present every half a minute</th> </tr> <tr> <th>0.0 min</th> <th>0.5 min</th> <th>1.0 min</th> <th>1.5 min</th> <th>2.0 min</th> <th>2.5 min</th> <th>3.0 min</th> <th>3.5 min</th> <th>4.0 min</th> </tr> </thead> <tbody> <tr> <td>Chloride ions present</td> <td>0</td> <td>24</td> <td>54</td> <td>70</td> <td>80</td> <td>88</td> <td>95</td> <td>100</td> <td>100</td> </tr> <tr> <td>Chloride ions absent</td> <td>0</td> <td>12</td> <td>20</td> <td>29</td> <td>36</td> <td>40</td> <td>45</td> <td>48</td> <td>50</td> </tr> <tr> <td>Difference in maltose concentration When chloride ions are either present or absent</td> <td>0</td> <td>12</td> <td>34</td> <td>41</td> <td>44</td> <td>48</td> <td>50</td> <td>52</td> <td>50</td> </tr> </tbody> </table> <p>Allow a + /- 1% for any concentration of maltose and a + /- 2% for the difference in maltose concentrations</p> | | Presence or absence of chloride ions | The percentage concentration of maltose (%) present every half a minute | | | | | | | | | | 0.0 min | 0.5 min | 1.0 min | 1.5 min | 2.0 min | 2.5 min | 3.0 min | 3.5 min | 4.0 min | Chloride ions present | 0 | 24 | 54 | 70 | 80 | 88 | 95 | 100 | 100 | Chloride ions absent | 0 | 12 | 20 | 29 | 36 | 40 | 45 | 48 | 50 | Difference in maltose concentration When chloride ions are either present or absent | 0 | 12 | 34 | 41 | 44 | 48 | 50 |
| Presence or absence of chloride ions | The percentage concentration of maltose (%) present every half a minute | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | 0.0 min | 0.5 min | 1.0 min | 1.5 min | 2.0 min | 2.5 min | 3.0 min | 3.5 min | 4.0 min | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Chloride ions present | 0 | 24 | 54 | 70 | 80 | 88 | 95 | 100 | 100 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Chloride ions absent | 0 | 12 | 20 | 29 | 36 | 40 | 45 | 48 | 50 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Difference in maltose concentration When chloride ions are either present or absent | 0 | 12 | 34 | 41 | 44 | 48 | 50 | 52 | 50 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | (c) (ii) | <p>1 (acts as a) cofactor ;</p> <p>2 (Cl⁻) binds to, enzyme / amylase / amylose / substrate ;</p> <p>3 enzyme substrate complex / ESC, forms more, easily / quickly ;</p> | 2 max | <p>1 IGNORE ‘coenzyme’</p> <p>2 ACCEPT binds to, active site</p> <p>3 ACCEPT description</p> | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

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| (c) | (iii) | <p>1 temperature ;</p> <p>2 pH ;</p> <p>3 enzyme / amylase / chloride, <u>concentration</u> ;</p> <p>4 substrate / starch / amylose, <u>concentration</u> ;</p> <p>5 constant / regular, stirring ;</p> <p>6 (fixed) <u>volume</u> of solution (removed each time for sampling) ;</p> | 3 max | <p>Mark the first three answers only regardless of which line they are on DO NOT CREDIT refs to, time</p> <p>3 IGNORE 'amount' or 'volume' 3 DO NOT CREDIT 'concentration' unqualified</p> <p>4 IGNORE 'amount' or 'volume' 4 DO NOT CREDIT 'concentration' unqualified</p> |
| | | Total | 19 | |

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|----------|-----------|--|------|---|
| 4 | (a) (i) | <p>1 (m)RNA is single stranded / DNA is double stranded ;</p> <p>2 (m)RNA is non helical / DNA is helical ;</p> | 1 | <p>Mark the first response but do not award the mark if a further answer is incorrect or contradictory DO NOT CREDIT refs to length as given in stem</p> <p>1 ACCEPT DNA is a double helix (as stranded is implied) for this mp DO NOT CREDIT DNA is a double <i>molecule</i></p> <p>2 ACCEPT (mRNA) not twisted / not coiled / not spiral / straight / ora</p> |
| | (a) (ii) | <p>1 RNA contains ribose and DNA contains deoxyribose ;</p> <p>2 RNA contains, uracil / U, and DNA contains, thymine / T ;</p> <p>3 3 / more than 1, forms of RNA ;</p> <p>4 RNA is, single stranded / non helical, and DNA is, double stranded / helical ; <i>if not already awarded as answer in (i)</i></p> | 1 | <p>Mark the first response to (a)(ii) – but but do not award the mark if a further answer is incorrect or contradictory</p> <p>2 DO NOT CREDIT thymine</p> <p>3 ACCEPT 'one form of DNA'</p> |
| | (a) (iii) | <u>gene</u> ; | 1 | IGNORE allele / operon |
| | (a) (iv) | too big to / does not, fit through <u>pore</u> (in nuclear envelope) ; | 1 | ACCEPT 'too long to fit ... pore' |
| | (a) (v) | <p><i>idea that</i> only copies one, gene / section / part / AW, (of DNA) ;</p> <p><i>idea that</i> DNA comprises many, genes / alleles ;</p> | 2 | <p>e.g. mRNA only codes for 1 protein</p> <p>DO NOT CREDIT '1 DNA molecule contains <u>all</u> the genes' 'mRNA only codes for 1 protein but DNA codes for many proteins' = 2 marks</p> |

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| (b) | (i) | <p>1 <u>non-competitive</u> (inhibitor) ;</p> <p>2 (α-amanitin / inhibitor / toxin) fits into, allosteric site / a place other than active site ;</p> <p>3 <u>active site</u> changes, shape / configuration / conformation / structure ;</p> <p>4 substrate no longer, fits / complementary to, <u>active site</u> ;</p> | 2 max | <p>3 ACCEPT 'distortion of active site'</p> <p>4 Mark to be awarded in context of active site (although need not be repeated if stated in mp 3) IGNORE ESC</p> |
| (b) | (ii) | <p>1 inhibits production of mRNA / mRNA not produced ;</p> <p>2 prevents protein synthesis / AW ;</p> <p>3 e.g. of, specific named protein / (vital) process, that may be affected ;</p> | 2 max | <p>1 CREDIT prevents transcription</p> <p>2 CREDIT translation</p> <p>3 e.g. respiration / photosynthesis (as question refers to 'an organism') / haemoglobin / cytochrome C oxidase</p> |
| (c) | (i) | sequence / order, of amino acids ; | 1 | IGNORE number / organisation |
| (c) | (ii) | <p>A = ionic ;</p> <p>B = hydrogen ;</p> <p>C = <u>disulfide</u> (bond / bridge) ;</p> | 3 | <p>ALLOW phonetic spelling</p> <p>DO NOT CREDIT <u>disulfate</u></p> |
| (d) | | <p>1 increased <u>kinetic</u> energy ;</p> <p>2 (any part of protein molecule) vibrates ;</p> <p>3 hydrophilic / hydrophobic / hydrogen / ionic, bonds / interactions, break ;</p> <p>4 change in, <u>3D</u> shape / conformation (of protein) ;</p> <p>5 <u>denatures</u> ;</p> | 3 max | <p>1 must contain the idea of <u>more</u> than normal</p> <p>3 IGNORE Van der Waals DO NOT CREDIT if disulfide / covalent / peptide bonds are included</p> <p>4 IGNORE tertiary / structure (as in question) IGNORE refs to, active site / enzymes</p> |
| Total | | | 17 | |