

## Proteins - Mark Scheme

Q1.

| Question Number | Answer  | Additional Guidance   | Mark       |
|-----------------|---|---|------------|
|                 | <p>An answer which makes reference to four of the following:</p> <ul style="list-style-type: none"> <li>• both are chains of amino acids joined by peptide bonds (1)</li> <li>• both contain named bonds (holding molecule in its three dimensional shape) (1)</li> <li>• globular proteins have hydrophilic groups on the outside whereas fibrous proteins have hydrophobic groups on the outside (1)</li> <li>• globular have tertiary or quaternary structures whereas fibrous have little or no tertiary structure (1)</li> <li>• globular are folded into compact shapes whereas fibrous have long chains (1)</li> </ul> | <p>i.e. hydrogen bonds, disulfide bridges, ionic bonds</p> <p>Allow converse</p> <p>ALLOW globular being spherical and fibrous being long strands</p> | <b>(4)</b> |

Q2.

| Question Number | Answer  | Additional Guidance  | Mark       |
|-----------------|---|--|------------|
|                 | <p>An answer that makes reference to the following:</p> <ul style="list-style-type: none"> <li>• (proteins which) reduce activation energy of biological reactions (1)</li> </ul> | <p>ALLOW increase rate of biological reactions</p> <p>ALLOW references to {in cells / in living organisms}</p> | <b>(1)</b> |

## Q3.

| Question Number | Answer  | Additional Guidance   | Mark    |
|-----------------|---|---|---------|
| (a)             | Diagram clearly showing:<br>1. central carbon with {R / H / eq} and H attached by single bonds ;<br>2. {NH <sub>2</sub> / NH <sub>3</sub> <sup>+</sup> } attached to carbon by single bond ;<br>3. {COOH / COO <sup>-</sup> } attached to carbon by single bond ; | 1. Must show C, H and R or a plausible R group<br>2. and 3 ACCEPT groups attached to a central C that is not shown (chemical notation)<br>ACCEPT groups written wrong way round e.g. C-H <sub>2</sub> N<br>NOT incorrect bonding within groups e.g. C=OH<br>ACCEPT if correct group attached to wrong molecule e.g. glucose | (3) Exp |

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|-----------------|--|--|---------|
| (b) (i)         | 1. idea that enzymes reduce activation energy ;<br>2. reference to active sites (of enzyme) ;<br>3. reference to effect on collisions between enzymes and substrates / enzyme substrate complexes / eq ;<br>4. idea of number of active sites occupied ;<br>5. (levels off when) substrate becomes limiting factor ; | IGNORE increases the rate of the reaction<br>1. Accept 'decreases energy needed for reaction', provides an alternative reaction pathway<br><br>4. ACCEPT below 6a.u. all sites occupied OR above 6 a.u. not all occupied | (3) Exp |

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|-----------------|--|---|---------|
| (b) (ii)        | 1. idea of a range of concentrations of enzyme (at least 5) ;<br>2. idea of substrate concentration not limiting ;<br>3. reference to mixing ;<br>4. description of how to measure dependent variable with time ;<br>5. description of how to measure the initial rate of reaction ;<br>6. reference to an appropriate named controlled variable ;<br>7. reference to {replicates / repeats} at each enzyme concentration ;<br>8. control {described / used as comparison} ; | 4. and 5. Must relate to reaction / enzyme named<br><br>5. ACCEPT clear indication of rate measured soon after mixing, plot and calculate rate from linear part of graph<br>NOT time taken for all substrate to be converted but could get Mp4<br><br>6. ACCEPT e.g. pH, temperature, volume, concentration of substrate<br><br>7. IGNORE repeat for other concentrations<br>ACCEPT repeat whole experiment<br><br>8. ACCEPT control used is with {no enzyme / distilled water} | (4) Exp |

Q4.

| Question Number | Acceptable Answer | Additional guidance | Mark |
|-----------------|-------------------|---------------------|------|
| (i)             | C                 |                     | (1)  |

| Question Number | Acceptable Answer  | Additional guidance | Mark |
|-----------------|--|---------------------|------|
| (ii)            | <p>water molecule indicated (1)</p> $\begin{array}{c} \text{H} & \text{O} & \text{H} \\ & \diagdown & / \\ & \text{O} & \end{array}$ <p>correct dipeptide shown (1)</p> $\begin{array}{ccccccc} \text{H} & & \text{H} & \text{O} & \text{H} & \text{H} & \text{O} \\ & &   &    &   &   &    \\ \text{H} & - & \text{N} & - & \text{C} & - & \text{C} & - & \text{N} & - & \text{C} & - & \text{C} & - & \text{O} \\ & &   & &   & &   & &   & &   & &   & & \text{OH} \\ & & \text{H} & & \text{H} & & \text{H} & & \text{H} & & \text{H} & & \text{H} & & \end{array}$ |                     | (2)  |

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|-----------------|-------------------|---------------------|------|
| (iii)           | C                 |                     | (1)  |

Q5.

| Question Number | Answer   | Additional Guidance  | Mark |
|-----------------|--|--|------|
|                 | <p>1. idea of formation of secondary or tertiary structure ;</p> <p>2. idea of bonding between R groups ;</p> <p>3. named bond e.g. ionic, disulfide, hydrogen ;</p> | <p>1. <b>ACCEPT</b> e.g. alpha helix, beta pleated sheet, globular structure<br/><b>ACCEPT</b> folding (of primary structure)<br/><b>IGNORE</b> 3D shape</p> <p>2. <b>ACCEPT</b> hydrophilic R groups go to outside/<br/>hydrophobic R groups go to inside / eq</p> <p>3. <b>DO NOT ACCEPT</b> peptide</p> | (3)  |

Q6.

| Question Number | Answer  | Additional Guidance  | Mark       |
|-----------------|---|--|------------|
|                 | <p>1. idea that this increased temperature changes the bonding in the enzyme ;</p> <p>2. the active site is {denatured / changes shape} ;</p> <p>3. the substrate no longer fits into the active site / the enzyme no longer {catalyses the reaction / lowers the activation energy / eq} ;</p> | <p><b>IGNORE</b> enzyme is denatured</p> <p><b>ACCEPT</b> bonds are broken</p> <p>3. <b>ACCEPT</b> no enzyme substrate complex can form / eq</p> | <b>(2)</b> |

Q7.

| Question Number | Answer  | Additional Guidance   | Mark       |
|-----------------|---|---|------------|
|                 | <p>An explanation that makes reference to the following:</p> <ul style="list-style-type: none"> <li>• (different primary structure) results in a different sequence of amino acids (1)</li> <li>• change in R groups changes { folding / bonding / secondary structure / tertiary structure } (1)</li> <li>• changing { shape / charge } of the active site prevents substrate from being able to bind (1)</li> <li>• { stopping / reducing } the production of fibrin (1)</li> </ul> | <p>Do not accept peptide bonds</p> <p>ALLOW another specific aspect of blood clotting cascade</p> | <b>(4)</b> |

Q8.

| Question Number | Answer  | Additional guidance   | Mark       |
|-----------------|---|---|------------|
|                 | <p>A description that makes reference to the following:</p> <ul style="list-style-type: none"> <li>• (a peptide bond is formed by a) condensation reaction (1)</li> <li>• between the {amine group / NH<sub>2</sub>} and the {carboxyl group / COOH} of adjacent amino acids (1)</li> </ul> | <p>ALLOW 'amino' for 'amine' and 'carboxylic acid' for 'carboxyl'</p> | <b>(2)</b> |