

- M1.(a)**
1. Dissolve in alcohol, then add water;
 2. White emulsion shows presence of lipid.
- 2
- (b) Glycerol.
- 1
- (c) Ester.
- 1
- (d) **Y** (no mark)
Contains double bond between (adjacent) carbon atoms in hydrocarbon chain.
- 1
- (e)
1. Divide mass of each lipid by total mass of all lipids (in that type of cell);
 2. Multiply answer by 100.
- 2
- (f) Red blood cells free in blood / not supported by other cells so cholesterol helps to maintain shape;
- Allow converse for cell from ileum – cell supported by others in endothelium so cholesterol has less effect on maintaining shape.*
- 1
- (g)
1. Cell unable to change shape;
 2. (Because) cell has a cell wall;
 3. (Wall is) rigid / made of peptidoglycan / murein.
- 2 max
- [10]**
- M2.(a)**
1. Starch formed from α -glucose but cellulose formed from β -glucose;

2. Position of hydrogen and hydroxyl groups on carbon atom 1 inverted.

2

- (b) 1. Insoluble;
2. Don't affect water potential;

OR

3. Helical;

Accept form spirals

4. Compact;

OR

5. Large molecule;
6. Cannot leave cell.

2

- (c) 1. Long and straight chains;
2. Become linked together by many hydrogen bonds to form fibrils;
3. Provide strength (to cell wall).

3

[7]

- M3.(a)** 1. Helicase;
2. Breaks hydrogen bonds;
3. Only one DNA strand acts as template;
4. RNA nucleotides attracted to exposed bases;
5. (Attraction) according to base pairing rule;
6. RNA polymerase joins (RNA) nucleotides together;
7. Pre-mRNA spliced to remove introns.

6 max

- (b) 1. Polymer of amino acids;
2. Joined by peptide bonds;
3. Formed by condensation;
4. Primary structure is order of amino acids;
5. Secondary structure is folding of polypeptide chain due to hydrogen bonding;

Accept alpha helix / pleated sheet

6. Tertiary structure is 3-D folding due to hydrogen bonding and ionic / disulfide bonds;
7. Quaternary structure is two or more polypeptide chains.

5 max

- (c) 1. Hydrolysis of peptide bonds;
 2. Endopeptidases break polypeptides into smaller peptide chains;
 3. Exopeptidases remove terminal amino acids;
 4. Dipeptidases hydrolyse / break down dipeptides into amino acids. 4 [15]
- M4.(a)** 1. Maltose;
 2. Salivary amylase breaks down starch. 2
- (b) Maltase. 1
- (c) (Mimics / reproduces) effect of stomach. 1
- (d) 1. Add boiled saliva;
 2. Everything same as experiment but salivary amylase denatured. 2
- (e) 1. Some starch already digested when chewing / in mouth;
 2. Faster digestion of chewed starch;
 3. Same amount of digestion without chewing at end.
Accept use of values from graph 3 [9]
- M5.(a)** 1. **A:** phospholipid (layer);
 1. *Reject hydrophobic / hydrophilic phospholipid*
 2. **B:** pore / channel / pump / carrier / transmembrane / intrinsic / transport protein;
 2. *Ignore unqualified reference to protein*

2

(b) (i) Condensation (reaction);

1

(ii) Organelle named; Function in protein production / secretion;

Function must be for organelle named

Incorrect organelle = 0

eg

1. Golgi (apparatus);

1. Accept smooth endoplasmic reticulum

2. Package / process proteins;

OR

3. Rough endoplasmic reticulum / ribosomes;

3. Accept alternative correct functions of rough endoplasmic reticulum. ER / RER is insufficient

3. Accept folding polypeptide / protein

4. Make polypeptide / protein / forming peptide bonds;

OR

5. Mitochondria;

6. Release of energy / make ATP;

6. Reject produce / make energy

6. Accept produce energy in the form of ATP

OR

7. Vesicles;

8. Secretion / transport of protein;

2

[5]