<b>M1.</b> (a)	<ol> <li>Dissolve in alcohol, then add water;</li> <li>White emulsion shows presence of lipid.</li> </ol>	2	
(b)	Glycerol.	1	
(c)	Ester.	1	
(d)	<b>Y</b> (no mark) Contains double bond between (adjacent) carbon atoms in hydrocarbon chain.	1	
(e)	<ol> <li>Divide mass of each lipid by total mass of all lipids (in that type of cell);</li> <li>Multiply answer by 100.</li> </ol>	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)	<ol> <li>Cell unable to change shape;</li> <li>(Because) cell has a cell wall;</li> <li>(Wall is) rigid / made of peptidoglycan / murein.</li> </ol>	2 max	[10]

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).

[7]

3

- 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 <u>and</u> ionic / disulfide bonds:
- 7. Quaternary structure is two or more polypeptide chains.

5 max

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

(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;

[5]