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Detailed mark scheme

Suitable for all boards

Designed to test your ability and thoroughly prepare you

2002

XVIII

1583

Time allowed

52 Minutes

Score

/43

Percentage

%

Biology

**AQA
AS & A LEVEL**

Mark Scheme

3.3 Organisms exchange substances with their environment

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1

(a) In one country where the percentage of fat (in the diet) is 35%, the death rate (from breast cancer) is 20 per 100 000;

Must have reference to country

Accept ... 1 per 5 000 / 0.02%

1

- (b)
1. No. of deaths from breast cancer divided by total population \times 100 000;
 2. No. of deaths from breast cancer divided by all deaths \times 100 000;
 3. Sample and count deaths from breast cancer in 100 000 people;
If sample not 100 000 then must scale appropriately

1 max

- (c)
1. Positive correlation;
 2. But correlation does not show causation / some other (named) factor may be involved;
 3. Evidence against positive correlation e.g. different death rates at same % fat / similar death rates at different % fat / some countries with higher death rate have lower fat intake;
1. Accept description of positive correlation / directly proportional.
Accept positive relationship.
2. Do not accept casual in place of causal.
3. Answer must be consistent with data.

3

[5]



- 2 (a) 1. Add iodine / potassium iodide solution to the food sample; 1.
Allow 'iodine'
2. *Must be in the context of the correct reagent*
2. Blue / black / purple indicates starch is present; 2
- (b) 1. Starch digested to maltose / by amylase;
Ignore 'hard to digest / easily digested'
2. Maltose digested to glucose / by maltase;
3. Digestion of sucrose is a single step / only one enzyme / sucrase;
3. *Accept converse for starch*
3. *Do not accept digestion of sucrose is faster* 3
- (c) 1. Smoking increases risk of CHD / introduces another variable; 1
- (d) (i) 1. No effect on risk with diet group 1 and 2 / lowest glycaemic load;
Simple statement of correlation is not enough for this mark
2. Above diet group 2 / in higher groups, risk increases as glycaemic load increases; 1 max
- (ii) 1. (Higher GL diets lead to) more (harmful) lipids (in blood), so greater risk of atheroma;
Ignore reference to lipids in diet
2. Atheroma leads to blockage of coronary artery / increased risk of blood clot in coronary artery;
Ignore references to myocardial infarction / heart attack 2

[9]

- 3 (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]**
- 4 (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]

5 (a) C.

Ignore name of organ

1

(b) E.

Ignore name of organ

1

- (c)
1. Active site (of enzyme) has (specific) shape / tertiary structure / active site complementary to substrate / maltose;
Reject active site on substrate.
Must have idea of shape
Assume "it" = maltase
Accept (specific) 3D active site
Reject has same shape
 2. (Only) maltose can bind / fit;
Accept "substrate" for "maltose"
 3. To form enzyme substrate complex.
Accept E-S complex

3

[5]