UNIVERSITY OF CAMBRIDGE INTERNATIONAL EXAMINATIONS

GCE Advanced Subsidiary Level and GCE Advanced Level

MARK SCHEME for the October/November 2010 question paper for the guidance of teachers

9700 BIOLOGY

9700/36

Paper 32 (Advanced Practical Skills 2), maximum raw mark 40

This mark scheme is published as an aid to teachers and candidates, to indicate the requirements of the examination. It shows the basis on which Examiners were instructed to award marks. It does not indicate the details of the discussions that took place at an Examiners' meeting before marking began, which would have considered the acceptability of alternative answers.

Mark schemes must be read in conjunction with the question papers and the report on the examination.

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| Page 2 | Mark Scheme: Teachers' version | Syllabus | Paper |
|--------|-------------------------------------|----------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| Quest | ion | Expected Answers | | | | Additional guidance |
|-------------------------|-------|---|---------------------|---------------------------------|--|---------------------|
| 1 (a | | Decide on the temperature in the space below. | s you plan to | inge (between) 25°C to 45°C. | Record the temperatures you have chosen [2] | |
| 3.2 | [1] | at least 5 temperatures; | | | | |
| MMO | [1] | one temp. 25°C to 29°C | AND one tel 45°C | mp 40°C to | AND any three with two even intervals 3 or more degrees; | |
| | (ii) | Prepare the space below a | nd record you | ır results. | | [4] |
| 2 | [1] | Reject if any units in body of table only t | | | | |
| PDO recording 2 | | table with all cells drawn | AND head temperatu | ling (top or left re °C; | i) | Must have units |
| PDO rec | [1] | Reject if units in body of table if headings for volumes or stages (heading) time with units; | | | | |
| IO tion 2 | [1] | temperatures recorded highest to lowest | | ND st set of times | recorded in whole seconds; | |
| MMO collection 2 | [1] | time at the lowest tempera | ture is greater | temperature; | Allow • only if 3 or more results | |
| | (iii) | From your results, state th | e temperature | e activity of the enzyme is lov | west. [1] | |
| ACE interpretation 1 | [1] | temperature with longest ti | me | AND with u | nits, °C; | |

| Page 3 | Mark Scheme: Teachers' version | Syllabus | Paper |
|--------|-------------------------------------|----------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| | (iv) I | dentify two significant sources of error i | n this investigation. | | [2] |
|--------------------------|--------|--|--|-------|-----|
| | | cause of error | error | | |
| max 2 | [1] | (dependent) stage 3 or end-point clots stick small clots coagulation milk drains back slowly | idea of seeing determining judging when; | | |
| ACE interpretation max 2 | [1] | (standardised variables) rotation or angle; | AND idea of not constant/different not same | | |
| ACE | [1] | shaking or mixing or E/enzyme starts to react; | timing delayed; | | |
| | [1] | E/enzyme temperature; (as milk)/AW | | | |
| | [1] | (independent variable) temperature or test-tube removed from water-bath | idea of not constant/not maintained decreasing cools down; | Max 2 | |
| | | Describe a suitable control for this inves Reject if give two. | tigation. | | [1] |
| ACE improvement | [1] | boil enzyme; | | | |

| Page 4 | Mark Scheme: Teachers' version | Syllabus | Paper |
|--------|-------------------------------------|----------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| | (vi) Sugge | st how you co | uld mak | e this inve | stigation a | as reliable | as possib | ole. | | [1] |
|---|---|---------------------------|-------------|---|-------------|---------------------------------|------------|-----------------------|---|------------|
| ents MAX 1 | C control of any relevant variable | Or use thermosta Or | atically co | d enzyme to temp. separately then mix Ily controlled water bath r bath during rotation; | | | | | | |
| ACE improvements MAX 1 | R1 improve method to get repeat data [1] | repeat | | AND calculate or find mean/average; | | | | | | |
| (| í (ìií) Compl | ete the Table | 1.1 by ca | lculating t | | | cle around | each of these values. | 1 | [1] [1] |
| | [1] | circles around | 1 8.2, 4.9 | <u>, 1.1;</u> | | | | | | |
| | | | | | | ilk clotting er itrary units | nzyme | | | |
| MMO decisions 1 ACE interpretation 1 | | pH of milk | trial 1 | trial 2 | trial 3 | trial 4 | trial 5 | mean | | |
| isio reta | | 6.02 | 8.8 | 8.7 | 8.9 | (8.2) | 8.7 | 8.8 8.7 | | |
| dec | | 6.22 | 6.8 | 6.8 | 6.8 | 6.7 | 6.9 | 6.8 | | |
| MO: | | 6.40 | (4.9) | 4.3 | 4.4 | 4.3 | 4.4 | 4.4 | | |
| ACE M | | 6.64 | 1.1 | 1.0 | 1.0 | 0.9 | 1.0 | 1.0 | | |
| | | 6.70 | 0.7 | 0.6 | (1.1) | 0.5 | 0.7 | 0.6 | | |
| | [1] | 8.8 Allow 8.7 | | | | | | | | |

| Page 5 | Mark Scheme: Teachers' version | Syllabus | Paper |
|--------|-------------------------------------|----------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| | (i | iii) Plot a graph of the data shown in Table 1.1 | - | [4] |
|------------|----------|--|---|--|
| | O [1] | x-axis pH | AND y-axis activity (/) arbitrary units or au; | Must have units |
| | S | Reject if awkward scale | | error carried forward if |
| | [1] | scale as 0.2 to 2 cm Origin must be labelled as 6 or 6.02 | AND 2 to 2 cm; | incorrect O then scale x-axis 2 to 2 cm and y-axis 0.2 to 2 cm. must use more than half grid in x and y. |
| ayout 4 | Р | Reject plotting if scale is awkward if only dots/blobs or blobs in circles | intersection of cross must be clear to show plot. | |
| 100 | [1] | correct plotting using crosses/dots in circle only; | | |
| PDO layout | L [1] | straight line through points; error carried forward if scale or plotting incorrect 6.02 8.8 or 8.7 or ecf 6.22 6.8 6.40 4.4 6.64 1.0 6.70 0.6 | quality – not thick, not feathery for the complete line. joining plots – • ruled lines plot to plot • line of best fit • curve through all plots | |

| Page 6 | Mark Scheme: Teachers' version | Syllabus | Paper |
|--------|-------------------------------------|----------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| | (iv) E | xplain the relationship between pH | and the enzyme shown in the data. | [3] |
|-------------------|--------|--|--|-----|
| ACE conclusions 3 | [1] | (in correct context of pH and effect on activity) structure of protein or substrate or enzyme or active site | changed/altered/destroyed/no longer complementary broken; | |
| | [1] | (in correct context of increase in pH | and decrease activity) ces (ESCs) or less/no substrate can crease in activity) | |
| | [1] | (in correct context of effect of pH or acidic/more alkaline) denatured/denaturation; | | |
| | I | | [Total: 20] | |

| Page 7 | Mark Scheme: Teachers' version | Syllabus | Paper |
|--------|-------------------------------------|----------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| Ques | stion | Ex | pected Answers | | Additional guidance | |
|--------------------|---------|--|---|-----------------------------------|------------------------|-----|
| 2 (| (a) (i) | Draw a large plan diagram showing | the features of th | ne wall of the organ. Label the | position of the lumen. | [4] |
| 1 | [1] | Reject if drawn over print of question | | | | |
| PDO layout 1 | | Reject | AND no shading | AND uses most of space provided; | | |
| n 2 | [1] | Reject if drawn two walls | | | | |
| collection | | no cells drawn | AND three layers drawn include any circles as only one layer; | | | |
| MMO | [1] | Reject if only two layers drawn innermost layer is wider than outerme | | | | |
| MMO decisions 1 | [1] | Reject If any label is biologically incorre Iabel within drawn area – e.g. be correct label with label line to or in lui | tween two walls | enging to other organs or plants. | | |

| Page 8 | Page 8 Mark Scheme: Teachers' version | | Paper | |
|--------|---------------------------------------|------|-------|--|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 | |

| | (ii) | Annotate (make note layers. | s with label lines) your | diagram to show one difference b | petween the outside layers and the inside [1] |
|---------------------|------------|--|---|---|--|
| max 1 | | | s of the diagram drawing. hth, unless have labelled on | diagram | |
| | | | outermost | innermost | |
| decision | [1] | thickness Reject cell wall | thin)ner) | think(er); | |
| ММО | [1] [1] | texture | smooth | rough; | |
| Σ | | cells/nuclei | Not clear/densely packed/ visible | Clear/less densely packed/(air) spaces/lots | |
| | [1] | Colours/staining of | Pink/red/grey/lighter/mo | | max 1 |
| (k |) (i) | Actual diameter of th largest nucleolus in c | | elled Y is 7.8 µm. Use this informa | tion to calculate the actual diameter of the [4] |
| MMO collection 2 | [1] | correct measurement of one nucleus, 11 to 15 mm; | | | Reject if no units |
| Colle | [1] | correct measurement of one nucleolus, 2 to 4.5 mm; | | Reject if no units | |
| PDO display 2 | [1] | (mean) adds three measurements AND shows division by 3; | | | |
| PE disp | [1] | answer to no more than 2 significant figures, (1 decimal place) between 1.1 and 6.4; | | Reject standard form | |

| Page 9 | Page 9 Mark Scheme: Teachers' version | | Paper |
|--------|---------------------------------------|------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| | (ii) | Suggest how you would make the | | [1] | | |
|--------------------|---|---|-------------------------|----------------------------------|--|-----|
| _ | [1] | different dimensions/diameters | | | | |
| ACE improvement 1 | | or use vernier callipers | | | | |
| E impro | | or (eyepiece) graticule | or (eyepiece) graticule | | | |
| AC | | or increase magnification or high poversolution; | | | | |
| | (iii) | Make a large drawing of the cell la | abelled X with thre | e complete cells touching cell X | | [5] |
| | [1] | Reject if drawn over print of question | | | | |
| PDO layout 1 | | Rejectthick linesfeathery lines2 'tails' or overlaps or gaps | AND | AND | | |
| | | clear, sharp, unbroken lines | no shading | uses most of space provided; | | |
| | [1] | only cell \boldsymbol{X} and three correct complet | | | | |
| ion 2 | [1] | nucleus with at least two distinct nuc | × × | | | |
| MMO collection 2 | | | | | | |
| MMO decisions 2 | [1] chromosomes drawn as two areas (no details of chromosomes shown); | | | | | |
| MN decisi | [1] | blue region/spindle around chromoso | | | | |

| Page 10 | Page 10 Mark Scheme: Teachers' version | | Paper |
|---------|--|------|-------|
| | GCE A LEVEL – October/November 2010 | 9700 | 36 |

| | (iv) | Prep | pare the space below so th | at it suitable for you to | compare the cells labelled X a | and Y. [5] |
|--------------------------|------------|--|----------------------------------|---|---|---|
| recording 2 | [1] | organise as a table or Venn diagram or ruled connected boxes | | headed (cell) <u>X</u> and (cell) <u>Y</u> | differences opposite each other; | X Y |
| PDO | [1] | head | ing for similarities/similarity/ | | | |
| MMO decision | [1] | has at least one correct similarity, cytoplasm or cell/plasma membrane or shape; | | | | |
| | [1] | Reject tick and cross without a key | | | if no organisation then mark points only if | |
| | | | feature | (cell) X | (cell) Y | in same sentence or following sentences. |
| ACE interpretation max 2 | | 1 | nucleus/nuclear membran | e absent/none/not clear | present/clear; | Allow two ticks for both present i.e. for cytoplasm and shape. |
| tion | [1] | 2 | nucleoli | absent/none/ | present/clear; | Sytopiasin and shape. |
| etai | [1] | 3 | cytoplasm | less/not granular | more/granular; | Allow differences even if not opposite |
| erpr | [1] | 4 | spindle fibres | present/visible | absent/none/not visible; | each other. |
| i | [1] [1] | 5 | chromosomes/chromatid(| s) present/visible | not visible; | |
| ÇE | | 6 | cytoskeleton | absent/not clear | present/clear/visible; | Allow difference on one side if e.g. use |
| 1 | [1] | 7 | cell size | small(er) | larg(er); | more or –er. |
| | | | | Similarities | | |
| | | | | | | max 2 |
| | | | | | [Total: 20] | |