

CAMBRIDGE INTERNATIONAL EXAMINATIONS

Cambridge International Advanced Subsidiary and Advanced Level

MARK SCHEME for the May/June 2015 series

9700 BIOLOGY

9700/52

Paper 5 (Planning, Analysis and Evaluation),
maximum raw mark 30

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 should be read in conjunction with the question paper and the Principal Examiner Report for Teachers.

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Page 2	Mark Scheme	Syllabus	Paper
	Cambridge International AS/A Level – May/June 2015	9700	52

Mark scheme abbreviations:

- ;** separates marking points
- /** alternatives answers for the same point
- R** reject
- A** accept (for answers correctly cued by the question, or extra guidance)
- AW** alternative wording (where responses vary more than usual)
- underline** actual word given must be used by candidate (grammatical variants accepted)
- max** indicates the maximum number of marks that can be given
- ora** or reverse argument
- ecf** error carried forward
- I** ignore
- mp** marking point (with relevant number)

Page 3	Mark Scheme	Syllabus	Paper
	Cambridge International AS/A Level – May/June 2015	9700	52

Question	Expected answer	Extra guidance	Mark
1 (a)	<p><i>independent</i>: <u>concentration</u> of amylase / enzyme ;</p> <p><i>dependent</i>: diameter / area of brown zone ;</p>	<p>I amount R (amylase) extract</p> <p>A radius of brown zone A starch free / digested starch / clear zone A area of brown zone minus well R blue zone</p>	[2]
(b)	<p>any 8 from:</p> <p><i>independent variable</i></p> <p>1 ref. to method of diluting the 0.5 g dm⁻³ amylase / stock amylase solution and to give a minimum of 5 dilutions ;</p> <p>2 ref. to concentrations from 0.5 g dm⁻³ downwards with correct units ;</p> <p>3 use of a <u>control</u> with example ;</p> <p><i>dependent variable</i></p> <p>4 ref. to a suitable method of measuring diameters / width / radius / area of brown zones ;</p>	<p>1 0.0 and 0.5 g dm⁻³ can be included in the number of dilutions A serial / series / simple / proportional / dilution as method OR a description. Use the formula $C_1V_1 = C_2V_2$ to make... A If the fungal extract is diluted instead of amylase but R mp5</p> <p>2 minimum of 3 other stated values between 0.5 g dm⁻³ and 0.0 g dm⁻³ must correspond to dilution method chosen ecf if no method given A 10 fold or a 50% reduction serial dilution</p> <p>3 A water / 0.0 g dm⁻³ / boiled or denatured extract / enzyme</p> <p>4 e.g. using (suitable) ruler / callipers / string and ruler R metre ruler I graticules OR use a (transparent) grid / graph paper and count the number of squares OR take a photograph and measure using one the methods for finding area NOTE: method of measuring must match what they have stated they are measuring.</p>	

Page 4	Mark Scheme	Syllabus	Paper
	Cambridge International AS/A Level – May/June 2015	9700	52

<p>5 ref. to testing the (fungal) extract/fungal amylase/fungal enzyme ;</p> <p>6 plot a calibration curve of known concentrations and use it to determine extract concentration ;</p> <p><i>standardising variables (max. 3)</i></p> <p>7 ref. to suitable stated volume/same volume of amylase (in each well) ;</p> <p>8 leave (all plates) for same period of time ;</p> <p>9 method of maintaining at same/constant/optimum/stated temperature ;</p> <p>10 use a buffer to keep the pH of the agar same ;</p> <p>11 same <u>concentration</u> of starch (in the agar plates) ;</p> <p>12 same depth/volume of agar in Petri dish ;</p> <p>13 cover to prevent contamination / evaporation ;</p> <p><i>safety</i></p> <p>14 ref. to low risk investigation/hazard <u>and</u> suitable safety precaution ;</p> <p><i>reliability</i></p> <p>15 ref. to a minimum of three replicates <u>and</u> calculate a mean or identify/eliminate/remove anomalies ;</p>	<p>5 R if dilute the fungal extract</p> <p>6 A compare extract result with results of known concentrations to identify extract concentration OR find approx. range of concentration and then do more at smaller intervals to identify extract concentration</p> <p>7 If a volume is stated – max 1 cm³ I amount/known</p> <p>8 if time stated, minimum 30 min/max 24 hours</p> <p>9 e.g. incubator/constant temp. room A water bath if temp. stated, any single temp. in range 15–65 °C I air conditioning</p> <p>10 e.g. making or adding buffer to agar or starch (solutions) A adding buffer to the amylase solution before using it A if stated, any single pH</p> <p>11 I ref. to other nutrients in agar, I amount</p> <p>12 A depth of agar plate I mass of agar/depth of wells</p> <p>14 fungal/enzyme allergy or fungus/enzyme/iodine/agar is irritant <u>and</u> wearing gloves/eye protection/mask I iodine as an allergen R no risk</p> <p>15 A original and 2 more/several/many/multiple A outliers for anomalies R reduce anomalies</p>	<p>[max 8]</p>
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Page 5	Mark Scheme	Syllabus	Paper
	Cambridge International AS/A Level – May/June 2015	9700	52

(c)	use a (glucose) biosensor / glucose dipstick or test-strip / named test strip, e.g. clinistix / uristix ; (because) γ -amylase will produce glucose only / β -amylase will produce maltose (mainly) ;	A use Benedict's test and EITHER weigh precipitate OR use colorimeter to find the <u>intensity of blue solution</u> left OR time the <u>first appearance</u> of a colour change A chromatography and stain (to show sugars) A Barfoed's test for monosaccharides <i>I idea that γ-amylase produces more glucose than β-amylase</i> R ora	[2]
(d) (i)	<u>1963</u> shown on extract B , plate 5 ; <u>1809</u> shown on extract D , plate 3 ;		[2]
(ii)	reject / eliminate / ignore / leave out (affected data from calculations) ;	A repeat until consistent results obtained I repeat unqualified OR repeat to find a mean	[1]

<p>(iii)</p>	<p>correct values for $\Sigma x = 1820$</p> <p>and $\bar{x} = 303 / 303.3 / 303.3^*$</p> <p>and $(x - \bar{x})^2$;</p> <table border="1" data-bbox="347 440 1220 576"> <tr> <td>$(x - \bar{x})^2$ allowed</td> <td>341</td> <td>341.3</td> <td>341.34</td> <td>341.4</td> <td>342</td> </tr> <tr> <td>$(x - \bar{x})^2$ not allowed</td> <td>341.32</td> <td>341.2</td> <td></td> <td></td> <td></td> </tr> </table> <p>correct value of s from table ;</p> <table border="1" data-bbox="347 676 1097 743"> <tr> <td>correct values of s (\pm)</td> <td>8.(0)</td> <td>8.26</td> <td>8.27</td> <td>8.3</td> </tr> </table>	$(x - \bar{x})^2$ allowed	341	341.3	341.34	341.4	342	$(x - \bar{x})^2$ not allowed	341.32	341.2				correct values of s (\pm)	8.(0)	8.26	8.27	8.3	<p>e.g.</p> <table border="1" data-bbox="1335 276 1742 389"> <tr> <td>Σ</td> <td>1820</td> <td></td> <td>342</td> </tr> <tr> <td>\bar{x}</td> <td>303</td> <td></td> <td></td> </tr> </table> <p>allow ecf for s from wrong value of $\Sigma (x - \bar{x})^2$</p>	Σ	1820		342	\bar{x}	303			<p>[max 2]</p>
$(x - \bar{x})^2$ allowed	341	341.3	341.34	341.4	342																							
$(x - \bar{x})^2$ not allowed	341.32	341.2																										
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Σ	1820		342																									
\bar{x}	303																											
<p>(iv)</p>	<p>any 3 from:</p> <p>1 the larger / AW the brown area the more amylase / more enzyme activity ora</p> <p>OR the larger / AW the brown area the more gene copies ; ora</p> <p>2 (person or extract) A had the highest / AW <u>concentration</u> of amylase / (person or extract) F has lowest AW <u>concentration</u> of amylase ;</p> <p>3 (person) A has the highest number of gene copies / (person) F has the lowest number of gene copies ;</p> <p>4 ref. to genetic variation in population (for the production of amylase) ;</p>	<p><i>Ignore ref. to proportionality / positive correlation between enzyme concentration and gene copies</i></p> <p>A strongest / largest / most / ora weakest / smallest / least</p> <p>A A>B>C>D>E>F</p> <p>I copying results without any conclusion about rank order</p>	<p>[3]</p>																									
<p style="text-align: right;">Total:</p>			<p>[20]</p>																									

Page 7	Mark Scheme	Syllabus	Paper
	Cambridge International AS/A Level – May/June 2015	9700	52

<p>2 (a)</p> <p>any 3 from: <i>variation in volunteers</i></p> <p>1 body mass / weight ;</p> <p>2 ref. to use of drugs e.g. medication / self-inflicted ;</p> <p>3 alcohol consumption ;</p> <p>4 smoking status ;</p> <p>5 ref. to ethnicity ;</p> <p>6 volunteer all have the same 'handedness' ;</p> <p>7 no medical / named medical condition affecting nerve conduction ;</p> <p><i>variation in method of applying test</i></p> <p>8 same arm tested ;</p> <p>9 idea of volunteer not moving (during the test) ;</p> <p>10 electrical charge / coulombs / voltage / potential difference ;</p> <p>11 same number of volunteers in each age category ;</p>	<p>Mark the first three given. <i>I environmental factors, e.g. temperature / light / noise in room / time of day</i> <i>I diet</i> <i>I ref. to distance between or position of electrodes / time of charge application</i></p> <p>7 e.g. Multiple Sclerosis(MS), Myalgic Encephalitis(ME), Muscular Dystrophy(MD), Motor Neurone Disease (MND), spinal bifida, polio A disease of the nervous system</p> <p>9 e.g. at rest / sitting / lying down</p> <p>10 A electrical stimulus I current</p>	<p>[max 3]</p>
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Page 8	Mark Scheme	Syllabus	Paper
	Cambridge International AS/A Level – May/June 2015	9700	52

(b)	age / years	mean conduction velocity $\pm S_M$	confidence limits		<i>note: must have both numbers correct for one mark</i>	[1]
			lower limit	upper limit		
	60–69	52.2 \pm 0.675	50.85	53.55 ;		
(c) (i)	30–39 and 70–79 OR 40–49 and 70–79 OR 50–59 and 70–79 ; there is no overlap of (confidence) limits / S_M ;				A descriptions, e.g. upper limit of 70–79 does not reach lower limit of 30–39 I error bars overlapping / range(bars) not overlapping I ref. to mean conduction velocity	[2]
(ii)	<i>test:</i> t test ; <i>reason:</i> comparing (two) <u>means</u> / normal distribution / continuous data ;				R continuous variation	[2]
(iii)	there is no <u>significant</u> difference between the (mean) conduction velocities / NCV of (individuals from) different age groups ;				<i>needs to be clear that the significant difference is in the conduction velocity and the not the ages</i> A the difference in (mean) conduction velocities of individuals from different age groups is not <u>significant</u> A stated age categories / women of different ages / young(er) and old(er) women / with age	1
(d)	large sample size / number of people tested ;				A tested 394 individuals / many people / lots of people ; I sufficient / enough people tested I number of different age categories / ref. to standard error	[1]
					Total:	[10]