Surname	Centre Number	Candidate Number
Other Names		2



GCE A LEVEL - NEW

A400U20-1





BIOLOGY – A level component 2 Continuity of Life

TUESDAY, 20 JUNE 2017 – MORNING 2 hours

For Examiner's use only		
Question	Maximum Mark	Mark Awarded
1.	15	
2.	18	
3.	17	
4.	17	
5.	11	
6.	13	
7.	9	
Total	100	

ADDITIONAL MATERIALS

In addition to this examination paper, you will need a calculator and a ruler.

INSTRUCTIONS TO CANDIDATES

Use black ink or black ball-point pen. Do not use gel pen. Do not use correction fluid.

Write your name, centre number and candidate number in the spaces at the top of this page.

Answer all questions.

Write your answers in the spaces provided in this booklet. If you run out of space, use the continuation pages at the back of the booklet, taking care to number the question(s) correctly.

INFORMATION FOR CANDIDATES

The number of marks is given in brackets at the end of each question or part-question.

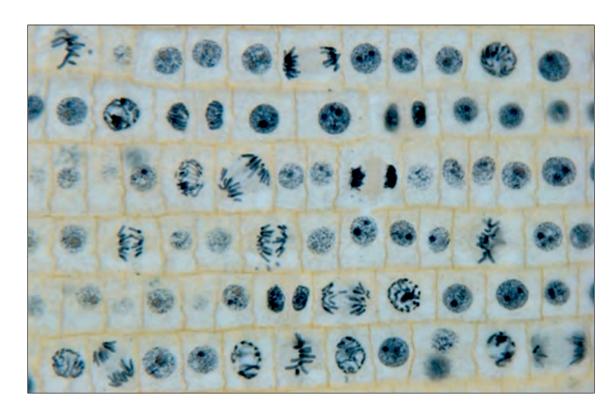
The assessment of the quality of extended repsonse (QER) will take place in question 7.

The quality of written communication will affect the awarding of marks.



Answer all questions.

1. The photomicrograph below shows cells from a root tip undergoing cell division.



(a) (i) Name the type of cell division taking place in this tissue. [1]

(ii) Count the number of cells in metaphase and anaphase. Enter the values in the table below: [2]

Phase	Number of cells	Time in minutes	% of cell cycle (by time)
Interphase	35	394	55
Prophase	9	101	14
Metaphase			
Anaphase			
Telophase/cytokinesis	10	112	16
Totals	64	720	100



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(iii)	Calculate the time taken for the two phases, using the formula.	
Time for a	а ра	$\frac{\text{number of cells in the phase}}{\text{total number of cells counted}} \times \text{total time taken for cell cycle}$	
		Enter the values in the table opposite.	[2]
(iv	')	Calculate the percentage time spent in each phase.	
		Enter the values in the table opposite.	[2]
(b) De	esc id a	ribe two ways in which the process of cell division shown differs between plant coanimal cells.	ells [2]
······	••••		
······			

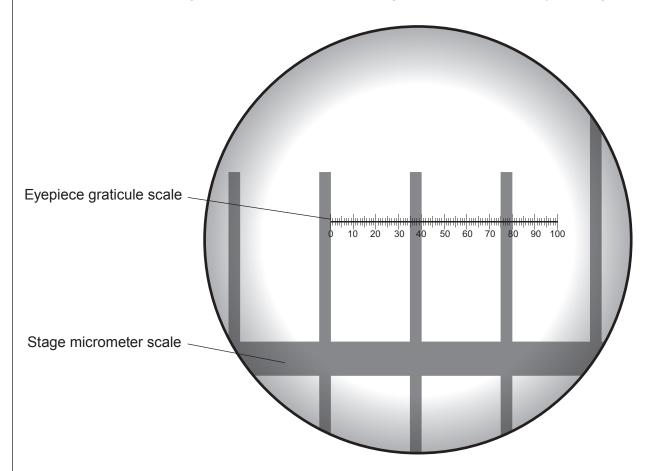


In hui tumoi		, unrestricted cell division can lead to the development of abnormal growths called		
(c)	(i)	What term is applied to the formation of malignant growths by unrestricted cell division? [1]		
	(ii)	Studies have shown that the plasma membranes of cells in these growths have some transmembrane proteins with shorter carbohydrate side chains.		
		In the boxes below, draw diagrams of a short section of the plasma membrane from a normal and an abnormal cell. Include labels for the following structures: phospholipid bilayer, transmembrane protein and carbohydrate side chains. [3]		
Normal		Abnormal		
	(iii)	How would the difference described in (ii) affect the properties of the abnormal cells?		





- 2. It is possible to germinate pollen grains *in vitro* by placing them in a suitable solution containing a mixture of mineral salts. One technique involves hanging a droplet of the mixture from the underside of a coverslip and suspending the coverslip above a microscope slide using petroleum jelly. The pollen grains may then be observed directly through a microscope.
 - (a) The photomicrograph below shows part of a stage micrometer and the eyepiece graticule.



The smallest divisions on the stage micrometer measure 100 µm.

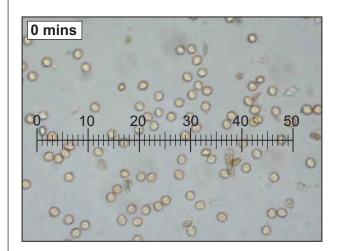
(i) Calibrate the eyepiece graticule. Express your answer to one decimal place. [2]

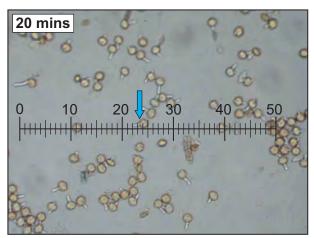


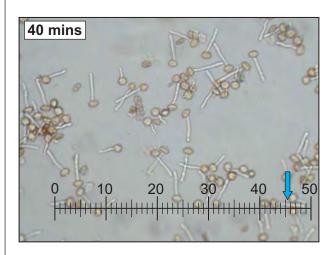
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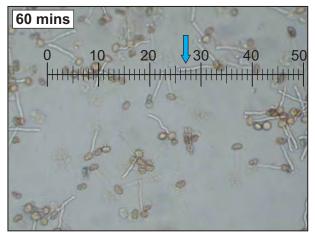
Turn over.

The photomicrographs below were taken at time intervals of 20 minutes from placing the pollen in the solution.





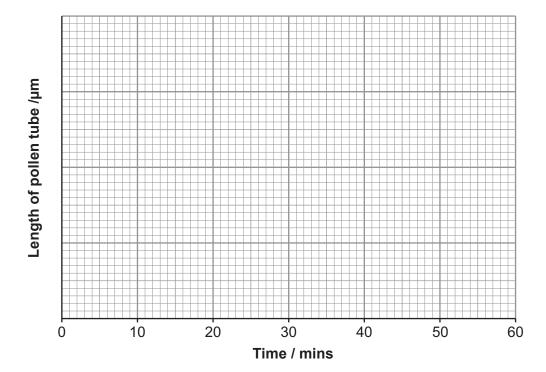




(ii) Use the eyepiece graticule to measure the length of the pollen tube indicated by an arrow from each photomicrograph. Measure each pollen tube to the nearest whole number. **Enter the values in the table below.** [2]

Time / mins	Length of pollen tube / eyepiece graticule units	Actual length of pollen tube /μm
0	0	0
20		
40		
60		

- (iii) Calculate the actual length of the pollen tubes in μm. **Write your answers in the table above**. [2]
- (iv) Plot a graph of the results using a suitable scale for the y-axis. [3]



(v) How could you make the data more reliable?

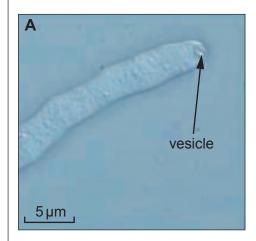
[1]

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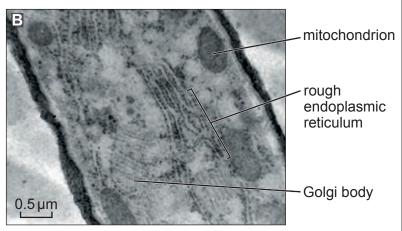
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The photomicrographs below show pollen tubes. Photomicrograph ${\bf A}$ shows a clear zone near the tip with an arrow; this is a vesicle. Electron micrograph ${\bf B}$ shows organelles present in a region further back from the tip.



(c)



tubes through the style of a flower.	[4]
	•••••••••••••••••••••••••••••••••••••••

Describe the roles of the vesicle and the labelled organelles in the passage of the pollen

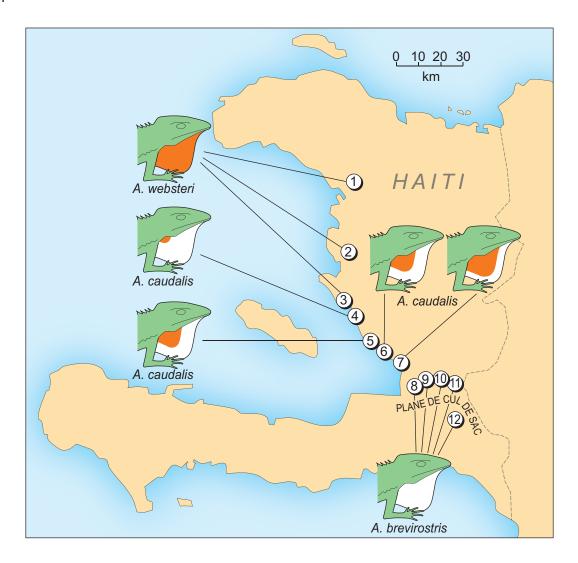
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3. In the Caribbean there is a group of lizards called Anoles. The males possess a retractable flap of skin under the throat, termed a dewlap, which is used in courtship, aggressive interactions and even encounters with predators. The patterns of orange colouration on the dewlaps of three species of Anole lizard and their distribution in Haiti are shown below.



Zone	Sites	Habitat
North	1,2,3	Beach
Central	4,5,6,7	Beach
South	8,9,10,11,12	Forest



0	
12	
5	
40	
- 2	~

(a)	(i)	Describe the patterns of orange and white dewlap colouration for each species.	
		[2	2]

Species	Pattern of orange and white
Anolis brevirostris	
Anolis websteri	
Anolis caudalis	
(ii) Describe	how the three species are distributed between the three zones. [1]
(iii) Describe	now the dewlap colouration varies from north to south in the central zone. [1]



Breeding experiments in the lab have revealed that dewlap colour-pattern is inherited as a Mendelian trait, in which orange (\mathbf{D}) is dominant to white (\mathbf{d}) .

The gene pool of the *A.brevirostris* population contains no **D** alleles.

The Hardy-Weinberg equations may be used to calculate the frequencies of the alleles $\bf D$ and $\bf d$ in the gene pool of the population of *A.websteri*

$$p + q = 1$$

 $p^2 + 2pq + q^2 = 1$

(b) In the north zone population of 400 lizards, four were found to have white dewlaps.

Calculate the frequencies of the two alleles in this population, and calculate how many of the lizards with orange dewlaps were heterozygous. [4]

1		Ш		Ш	
•		1	2		

The evolution of the three species is believed to be an example of sympatric speciation, resulting

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c)	(i)	Explain how the habitat in the south zone could bring about the dewlap colouration found in <i>A.brevirostris</i> .

	•••••	
	•••••	
	•••••	
	(ii)	Explain why after many generations the frequency of allele D is zero in <i>A.brevirostris</i> population but the frequency of allele d in <i>A.websteri</i> will never
	•••••	zero.
		zero.
d)	The	
d)	The	Anole lizards found at sites 3 and 4 are different species. Explain how their dew
d)	The colo	Anole lizards found at sites 3 and 4 are different species. Explain how their dewl

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Locus I.

This gene determines the colour of pigment. There are two major alleles: ${\bf B}$ coding for black pigment and ${\bf b}$ for brown.

Locus II.

This gene determines the distribution of pigment in the hair. Allele **A** produces a phenotype called agouti. The agouti phenotype is due to yellow bands on the otherwise dark hair shaft. In the non-agouti phenotype (determined by the allele **a**), the yellow bands are absent, so the hair shaft is a solid dark colour.

The table below shows the genotypes and corresponding phenotypes.

Locus I genotype	Locus II genotype	Phenotype (coat colour)
BB or Bb	AA or Aa	Normal agouti
BB or Bb	aa	Solid black
bb	AA or Aa	Cinnamon (brown with yellow bands)
bb	aa	Solid brown

bb		b	AA o	r Aa	Cinnamon	(brown with yellow	/ bands)
bb			a	a		Solid brown	
(a)	(i)		ne genetic di zygous at bo		w to show the	e expected outcon	ne of crossing
		Parental ph	enotype		x		
		Parental ge	notype		x		
		Gametes			x		
	(ii)	Give the rat	io in which th	e following p	ohenotypes w	ould appear in the	offspring. [1]
		Offspring pl	nenotypes:	normal ag	outi: solid bl	ack : cinnamon :	solid brown
		Phenotype	ratio:		:	:	

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A series of crosses between mice which were heterozygous at both loci produced 32 offspring. The results are shown in the table below. A null hypothesis was proposed that there was 'no significant difference between the observed and expected numbers of offspring'.

Phenotype	Observed O	Expected <i>E</i>	
Normal Agouti	22		
Solid black	4		
Cinnamon	3		
Solid brown	3		

(b) (i) Using the table provided, calculate χ^2 to test this null hypothesis. [3]

$$\chi^2 = \sum \frac{(O - E)^2}{E}$$

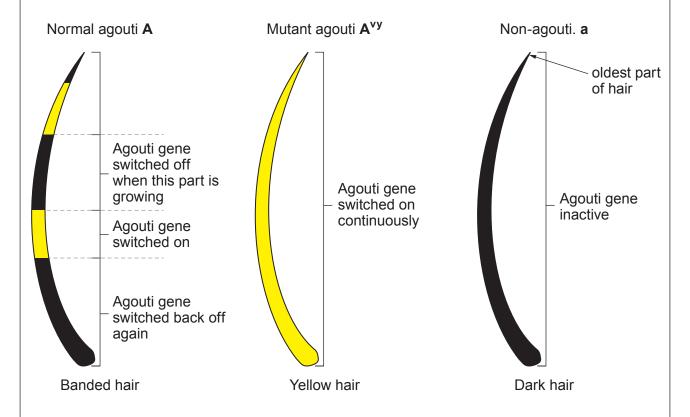
$$\chi^2 =$$

(ii) Choose an appropriate probability level and circle the critical value for χ^2 in the table below: [1]

Degrees	Probability								
freedom	0.9	0.8	0.7	0.5	0.2	0.1	0.05	0.02	0.01
1	0.016	0.064	0.15	0.46	1.64	2.71	3.84	5.41	6.64
2	0.21	0.45	0.71	1.39	3.22	4.60	5.99	7.82	9.21
3	0.58	1.00	1.42	2.37	4.64	6.25	7.82	9.84	11.34
4	1.06	1.65	2.20	3.36	5.99	7.78	9.49	11.67	13.28

(iii) State whether you accept or reject the null hypothesis, and explain why. [2]

A mutant form of the gene at locus II exists, called \mathbf{A}^{vy} , which results in hair which is yellow along its entire length. The diagrams of hairs below explain how each of the locus II alleles works.



Scientists have bred a strain of mice with the genotype $\mathbf{A}^{\mathbf{v}\mathbf{y}}\mathbf{a}$.

Since $\mathbf{A}^{\mathbf{v}\mathbf{y}}$ is dominant to \mathbf{a} , all of the mice with genotype $\mathbf{A}^{\mathbf{v}\mathbf{y}}\mathbf{a}$ could be expected to be yellow. In fact, some are yellow, some are normal agouti and some are shades in between.



These mice have identical DNA base sequences in the agouti gene locus but the gene is being expressed differently.

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(c) (i) What name is given to the study of the control of gene expression by factors other than changes in the DNA sequence? [1]

Normal allele (A)

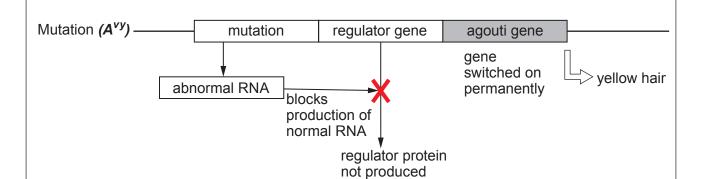
regulator gene

x

gene
switched on and off cyclically

regulator protein

regulator gene



Allele \boldsymbol{A} codes for the synthesis of a protein which switches off the gene in a cyclic manner

The mutation in A^{vy} is an insertion of a sequence of bases which forms abnormal RNA molecules which block the formation of the normal RNA.

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The addition of methyl groups to DNA molecules (methylation) is known to interfere with
the ability of the molecules to take part in the formation of mRNA.
It has been suggested that methylation of the inserted sequence of bases could be
responsible for the range of colours seen in the mice with genotype A ^{vy} a

sequence of bases results in mice with a range of coat colours. [3

17



used Fish to be	ently, a million tons of wild fish are harvested from the oceans annually. Of these, 80% are to feed fish reared in farms. do not make their own omega-3 fatty acids, they get these by eating algae. For farmed fish as nutritious as wild fish, they need to be fed a diet rich in omega-3 fatty acids. Algae are to culture on a large scale, so wild fish provide a more convenient source.
of on	ntists have identified and spliced genes from algae that code for producing high levels nega-3 fatty acids into <i>Camelina</i> plants. The genetically modified <i>Camelina</i> plants now uce seeds that contain 26% omega-3 fatty acids, so are useful for fish farm feed.
(a)	Explain how named enzymes would be used to cut genes from algae and splice them into Camelina plants. [3]
······	
(b)	Explain why feeding genetically modified <i>Camelina</i> seed to farmed fish could benefit biodiversity. [1]

Nutritionists often recommend that fish are included in diets, due to their high omega-3 polyunsaturated fatty acid content.

The diagrams show two omega-3 polyunsaturated fatty acids.

eicosapentaenoic acid (EPA)

$$\begin{array}{c} H \\ - O \\ O \\ \end{array}$$

docosahexaenoic acid (DHA)

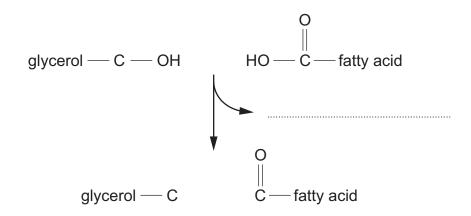
(c) With reference to the diagrams, explain why these compounds are described as:

(i)	fatty acids;	[2]
•••••		
(ii)	polyunsaturated.	[1]
•••••		

(d) (i) These compounds are present in fish in the form of triglygerides or phospholipids.

The diagram below shows some of the atoms of a glycerol molecule and a fatty acid molecule.

Complete the diagram to show how a bond is formed between the glycerol molecule and the fatty acid molecule, accounting for all of the atoms. [2]

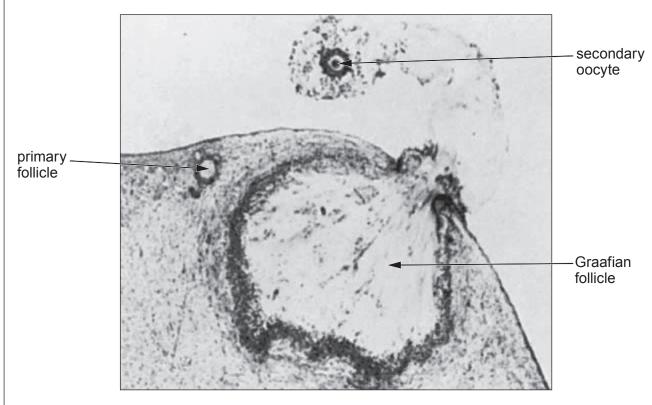


(ii) Give one function for the following types of lipid in fish: [2]triglyceride;

phospholipids.

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6. The photomicrograph below shows part of an ovary during ovulation.



(a) Describe what subsequently happens to the following: [2] secondary oocyte;

Graafian follicle.

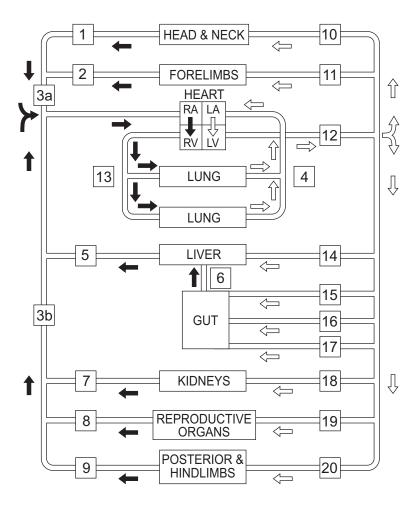
gative [4]	Examine only
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[3]	
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(b)	(i)	Describe how the hormones FSH and oestrogen illustrate the principle of neg feedback in controlling ovulation.	ative [4]
	<u></u>		
	·····		
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	(ii)	Explain why oestrogen may therefore be used as a contraceptive.	[3
	•••••		
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(c) For use as a contraceptive, oestrogen can be incorporated into a porous patch which may be applied to the skin on a woman's arm. The diagram below is a schematic map of the human circulatory system.



oxygenated blood ⇒ deoxygenated blood →



(i)	Using numbers for blood vessels and letters for the chambers of heart, describe the route that oestrogen would take from the patch to the organ where it would exert its contraceptive effect. [2]	Examiner only
(ii)	Apart from containing more oestrogen, describe one other difference in the blood leaving the patch and arriving at the target organ and state where the change would have taken place. [2]	
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7. The deep regions of the oceans (1000 – 4000 m) are nutrient-poor habitats. In order to survive, the fish that live there have evolved remarkable morphological and life-cycle strategies. The larvae of many deep-sea species develop in the nutrient-rich surface waters (0–200 m). Very few specimens of these fish have been caught, which makes them difficult to classify. This illustrates the tentative nature of classification. The table below provides information on specimens of three types of deep-sea fish.

Type of Fish	Example specimen	Evidence
Tapetail		Depth Caught: 0 – 200 m Number caught: 120 Size: 5 – 56 mm Age: juvenile Gender: unknown Feeding: large numbers of animal plankton; mouth is upturned, jaws are small
Whalefish		Depth Caught: below 1000 m Number caught: 600 Size: 26 – 408 mm Age: adult Gender: all female Feeding: eat large prey; jaws are long and can open very wide to swallow prey whole
Bignose Fish		Depth Caught: below 1000 m Number caught: 65 Size: 34 – 68 mm Age: adult Gender: all male Feeding: do not feed; have a very large liver which acts as a store of energy and nutrients



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Analysis of ribosomal RNA from examples which have been caught has produced the following phylogenetic tree. Phylogenetic evidence Procetichthys kreffti (Whalefish) Parataeniophorus gulosis (Tapetail) Cetostoma regain (Whalefish) Danacetichthys galathenus (Whalefish) Ataxolepis apus (Bignose fish) Eutaeniophorus festivus (Whalefish) Two hypotheses have been proposed to classify these deep-sea fish: Tapetails, Whalefish and Bignose Fish belong to different families of fish. **Hypothesis 1:** Tapetails, Whalefish and Bignose Fish are developmental stages of different **Hypothesis 2:** species of fish within the same family. For each type of evidence listed opposite and above, state whether they provide support for hypothesis 1 or hypothesis 2 and give reasons for your answers. From this evidence, conclude which hypothesis is more likely and suggest why a different conclusion may be reached in the future.



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