SurnameCentre
NumberCandidate
NumberOther Names2



GCE A level

1075/01

BIOLOGY/HUMAN BIOLOGY – BY5

A.M. FRIDAY, 20 June 2014

1 hour 45 minutes

For Examiner's use only							
Question	Maximum Mark	Mark Awarded					
1.	8						
2.	6						
3.	8						
4.	11						
5.	12						
6.	12						
7.	13						
8.	10						
Total	80						

INSTRUCTIONS TO CANDIDATES

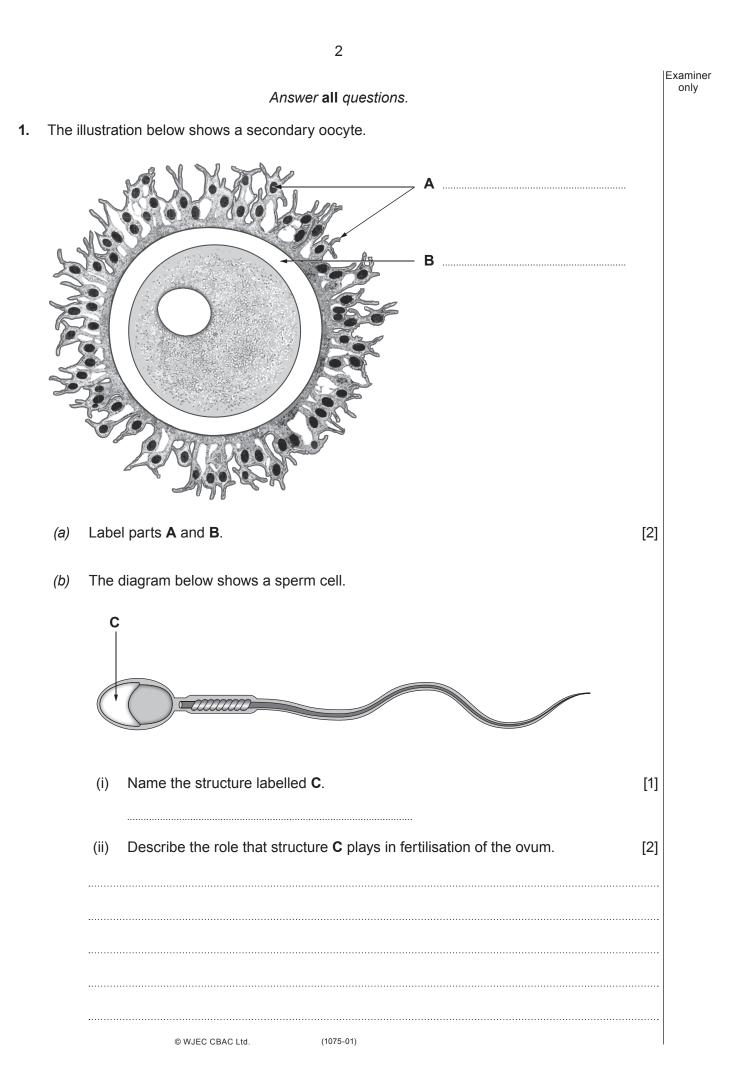
Use black ink or black ball-point pen.

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.

INFORMATION FOR CANDIDATES

The number of marks is given in brackets at the end of each question or part-question. You are reminded of the necessity for good English and orderly presentation in your answers. The quality of written communication will affect the awarding of marks. PMT



(C)	Expl	ain each of the following. [3]	Examiner only
	(i)	cell cleavage	
	••••••		
	••••••		
	(ii)	blastocyst	
	••••••		
	(iii)	implantation	
	•••••		ę
	••••••		1075 010003

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Examiner only The diagram below illustrates replication of DNA in cells. 2. G G l G G Describe the sequence of events shown within the dotted rectangle in the diagram (i) (a) above. [3] (ii) What is the role of DNA polymerase in the process? [1]

(b)	Explain why the process is referred to as 'semi conservative'. [2]	Examiner only
•••••		

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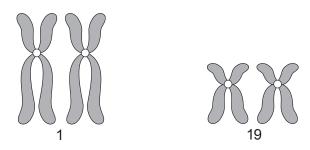
Examiner

only

[1]

3. The photograph below shows the pairs of chromosomes found in a body cell of a mouse.

- (a) What is the diploid number of the mouse?
- (b) The chromosomes in pairs 1 and 19 are commonly represented diagrammatically as:

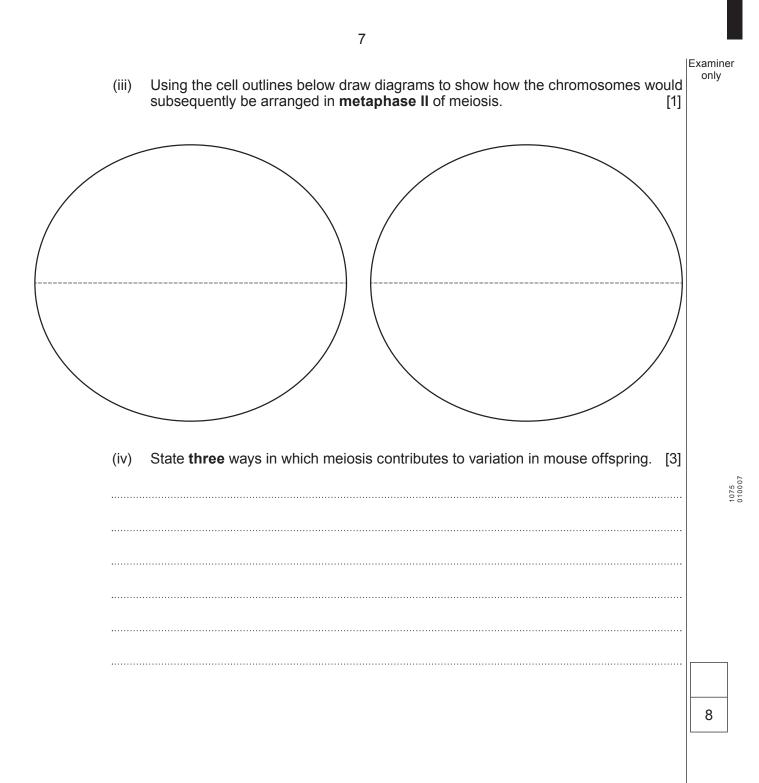


(i) Using the cell outline below draw diagrams to show how these pairs of chromosomes are arranged in **metaphase I** of meiosis. [1]

equator

(ii) On your drawing label; chromatid, centromere, centriole, spindle fibres.

[2]



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4. The fruit fly *Drosophila melanogaster* is extensively used to study genetics because it is relatively easy to cause mutations in the flies. Some mutant flies have very small (vestigial) wings:



normal wings



vestigial wings

Other mutants have very dark (ebony) bodies instead of the normal grey body.



grey body



ebony body

In a **dihybrid** cross, when flies with normal wings and grey bodies were crossed with flies with vestigial wings and ebony bodies all the offspring had normal wings and grey bodies.

(a) The F₁ hybrid flies (heterozygous for both traits) were allowed to interbreed freely. The F₂ flies were sorted and counted. The results are shown below.

Phen	Number of flies		
Wings	Body	Number of mes	
Normal	Grey	75	
Normal	Ebony	23	
Vestigial	Grey	21	
Vestigial	Ebony	9	

(i)	Draw a ge F ₂ phenotyp Use the lett	pe ratio.	ace provided	below, to show the expected [5]	Examiner only
	Allele for no	ormal wings = N	Allele for ves	stigial wings = n	
	Allele for gr	ey body = G	Allele for ebo	ony body = g	
F ₁ phenotyp	Des	Normal wing, grey body	Х	Normal wing, grey body	
F ₁ genotype	es		. Х		
Gametes			. X		

F₂ phenotype **ratio**

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(ii) Using the F_2 phenotype ratio from part (i) calculate the **expected** number of each phenotype in the F_2 generation from a total of 128 offspring, and enter the values in the table below. [1]

Phenotype		Observed number (O)	Expected number (E)	(O – E)	(O – E) ²	$\frac{(O - E)^2}{E}$
Normal wings	Grey body	75				
Normal wings	Ebony body	23				
Vestigial wings	Grey body	21				
Vestigial wings	Ebony body	9				

- (b) Complete the other columns in the table and carry out a Chi square test, testing the Null Hypothesis that there is no significant difference between the observed and expected results.
 - (i) Use the last column in the table to calculate χ^2 .

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

- $\chi^{2} =$
- (ii) Use the 5% probability level and the correct number of degrees of freedom to **circle** the critical value of χ^2 in the table below. [1]

Degrees	Probability								
of 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

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(iii) State whether you would accept or reject the Null Hypothesis, for this cross and explain why. [1]

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[1]

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(c) In another cross, flies with ebony bodies and scarlet eyes were crossed with flies homozygous for grey body and red eyes. All the F_1 flies had grey bodies and red eyes. When the F_1 hybrid flies were crossed the following results were obtained:

Pher	- Number of flies		
Eyes	Body		
Red	Grey	91	
Red	Ebony	3	
Scarlet	Grey	2	
Scarlet	Ebony	32	

The table shows that some of the offspring were far more common than expected and some phenotypes were very rare. Explain both of these observations. [2]

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5.	The techniques of recombinant DNA technology and micro-propagation are used to produce	
	Genetically Modified Crops. The following summary is adapted from an account given on the	
	Food Standards Agency's web site [www.food.gov.uk]	

- 1. A plant with the desired characteristic is identified e.g. resistance to the herbicide 'Roundup'.
- 2. The specific gene that produces this characteristic is found in the plant's DNA and cut out.
- 3. To get the gene into the cells of the plant being modified, the gene needs to be attached to a carrier. A piece of bacterial DNA called a plasmid is joined to the gene to act as the carrier.
- 4. Once the gene is attached to the plasmid, a marker gene is also added to identify which plant cells take up the new gene.
- 5. The 'gene package' is put in a bacterium, which multiplies, to create many copies of the 'gene package'.
- A copy of the 'gene package' is dried onto a gold or tungsten particle and fired into a piece of tissue from the plant being modified. The particle carries the 'gene package' into the plant's cells.
- 7. The plant tissue is put into a selective growth medium so that only modified tissue develops into plants.
- (a) Explain how different types of enzymes are used in stages 2 and 3 to produce the 'gene package'.
 [4]

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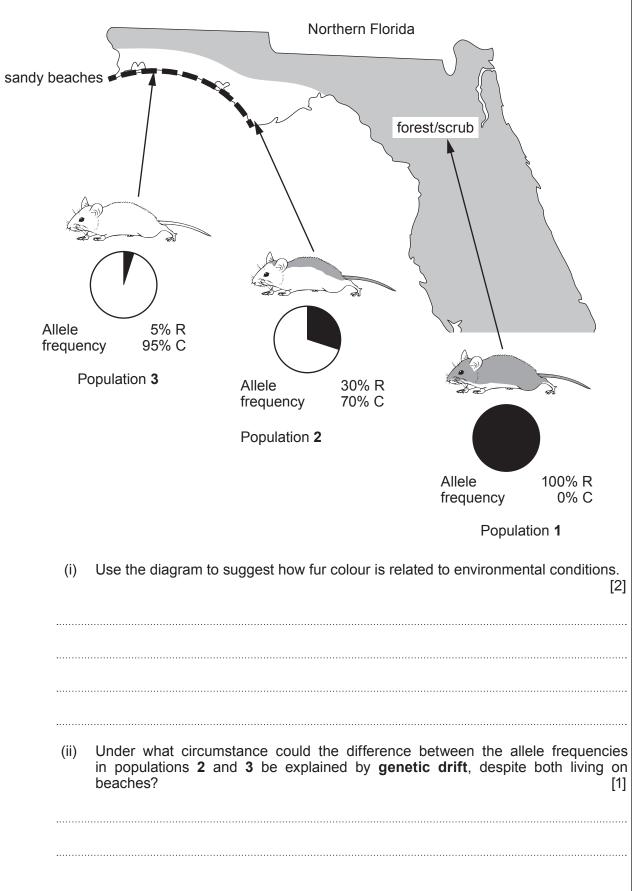
Examiner

- 6. A species of mouse *Peromyscus polionotus* found in Florida, USA, has a number of different coat colours. Coat colour in mice is controlled by several genes. Dark fur is produced when the hair producing cells secrete a pigment called eumelanin. A high level of eumelanin is produced when a transmembrane protein called MC1R is stimulated by a hormone.
 - (a) The diagram below shows part of the amino acid sequence of MC1R, part of the sequence of nucleotides in the gene for MC1R and how it might change to produce light fur:

Original		
Amino ac	id sequence	lle Thr Lys Asn Arg Asn Leu His Ser
Nucleotid (allele R)	le sequence	ATCACCAAAAACCGCAACCTGCACTCG
Changed	l to produce light	
Amino ac	id sequence	lle Thr Lys Asn Cys Asn Leu His Ser
Nucleotid (allele C)	le sequence	ATCACCAAAAACTGCAACCTGCACTCG
(i)	Describe the cha molecule.	ange in the gene and the subsequent change in the MC1R [2]
(ii)	Using the informa fur.	tion provided, explain how this change results in mice with light [2]

Examiner

(b) This change in the MC1R gene means that there are two alleles, R and C. The map below shows the distribution of the different coloured mice and the relative frequencies of the alleles R and C in each population.



	(iii)	Explain how Natural Selection could have caused the relative allele frequency shown in population 3 . [4]	Examiner only
	······		
	·····		
	(iv)	Under what circumstances would the mouse population become a separate species? [1]	
The	followi	ng is a quotation from an ecological investigation.	12
"Low unde	land h r threa	eaths are high-profile ecosystems for conservation action in England, but they are at from invasion by <i>Betula spp.</i> , <i>Pinus sylvestris</i> , and <i>Ulex europaeus</i> ."	
		[R.J. Mitchel et al. Journal of Applied Ecology, 1997, 37, 1426-1444]	
(a)	Disti	nguish between primary succession and secondary succession. [2]	

7.

Examiner only

The authors studied a number of heathland sites in Dorset including Arne, Blackhill, and Higher Hyde, where succession to one or another of the three species had taken place. The data below are based on the paper but have been simplified and modified for illustrative purposes. The successional stages in the study were named according to the dominant invasive species; **plus B**, where *Betula spp*, was the invader, **plus PS**, where *Pinus sylvestris* was the invader and **plus U**, where *Ulex europaeus*, was the invader.

(b) The group examined changes in soil chemistry from the original heath stage. Some of their results are summarised in the table below:

soil chemical property value by succession stage					
	original heath	plus B	plus PS	plus U	
рН					
Arne	3.63	4.01	3.60	3.63	
Blackhill	3.52	3.66	3.48	3.54	
Higher Hyde	3.53	5.06	3.51	3.47	
mean	3.56	4.24	3.53	3.55	
phosphorus µgPg ⁻¹					
Arne	2.41	3.85	2.69	3.16	
Blackhill	4.15	4.91	3.79	4.55	
Higher Hyde	5.08	5.35	3.55	4.76	
mean	3.88	4.70	3.34	4.16	
nitrate/nitrite µgNg ⁻¹					
Arne	0.51	0.65	0.59	1.16	
Blackhill	0.84	0.88	0.97	2.31	
Higher Hyde	0.69	0.98	1.17	3.64	
mean	0.68	0.84	0.91	2.37	

(i) What do the pH values tell us about the soil in all stages in all sites?

[1]

Use mean values from the table above to compare three changes to soil chemistry following invasion by *Betula spp*. with the changes following invasion by *Ulex europaeus*.

рН	 	
phosphorus		
nitrate/nitrite	 	

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% cover of species (by site) Species (by successional stage) Arne Blackhill Higher Hyde original heath Calluna vulgaris 62.0 66.1 88.2 Erica cinerea 22.4 25.7 2.6 Erica tetralix 9.9 2.6 9.9 0 Cladonia portentosa 8.5 0.5 plus B Betula spp. 18.9 11.7 16.5 Agrostis curtisii 0.0 53.6 0.0 Pteridium aquilinum 25.2 7.5 1.6 Calluna vulgaris 0.0 0.0 0.4 plus PS Pinus sylvestris 36.2 38.2 Pteridium aquilinum 0.3 24.7 Erica cinerea 0.0 0.0 Calluna vulgaris 0.0 0.0 plus U Ulex europaeus 87.0 75.3 79.0 Calluna vulgaris 14.7 5.8 7.2 Erica cinerea 1.5 11.3 4.3 Erica tetralix 0.1 0.3 0.3

(c) The table below shows changes to the vegetation in the successional stages:

(i) Which invading species has least impact on the vegetation on the original heathland? [1]

(ii) With reference to the data for **plus B** in both tables suggest a mechanism by which changes to vegetation occur during succession. [2]

		22	
(d)	Sixte (i)	een years later some of these successions have reached their natural conclusions. What name is given to the group of organisms that inhabit the ecosystem at the end of successional change?	Examiner only
	 (ii)	What usually happens to species diversity as succession proceeds? [1]	
	(iii)	Using named species from the table in part <i>(c)</i> explain why conservationists in Dorset are taking steps to prevent plus B and plus PS succession in heathland, but are less worried about type plus U succession. [2]	

8.				Examiner only
	Either,	(a)	Describe how the structure of a typical flower is adapted for insect pollination and subsequent fertilisation. [10]	
	Or	(b)	Describe energy transfer in an ecosystem. Briefly explain the agricultural practice of keeping animals in heated sheds with little room to move about. [10]	
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