



**General Certificate of Education (A-level)  
June 2013**

**Biology**

**BIOL2**

**(Specification 2410)**

**Unit 2: The Variety of Living Organisms**

**Final**

***Mark Scheme***

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| Question  | Marking Guidance   | Mark  | Comments   |
|-----------|--|-------|--|
| 1(a)(i)   | Centromere;  | 1     | Accept: if phonetically correct<br>Reject: centriole   |
| 1(a)(ii)  | <ol style="list-style-type: none"> <li>Holds chromatids together;</li> <li>Attaches (chromatids) to spindle;</li> <li>(Allows) chromatids to be separated/move to (opposite) poles / (centromere) divides/splits at metaphase/anaphase;</li> </ol> | 2 max | <ol style="list-style-type: none"> <li><b>Q</b> Neutral: chromosomes or chromatids split/halved/divided</li> <li>Reject: reference to homologous chromosomes being separated</li> </ol> Accept 'chromosomes' instead of 'chromatids'<br>Ignore incorrect names for <b>X</b>  |
| 1(a)(iii) | (Homologous chromosomes) carry different alleles;  | 1     | Accept alternative descriptions for 'alleles' eg different forms of a gene / different base sequences<br>Neutral: reference to maternal and paternal chromosomes   |
| 1(b)(i)   | (In <b>Figure 2</b> ) <ol style="list-style-type: none"> <li>Chromatids have separated (during anaphase);</li> <li>Chromatids have not replicated;</li> <li>Chromosomes formed from only one chromatid;</li> </ol>                                 | 1 max | <ol style="list-style-type: none"> <li><b>Q</b> Neutral: split/halved/divided</li> <li>Reject: reference to homologous chromosomes being separated</li> </ol> 1. & 2. Accept 'chromosomes' instead of 'chromatids'<br>Accept converse arguments for <b>Figure 1</b><br>Ignore references to the <i>cell</i> not dividing as in the question stem<br>Ignore: named phases |
| 1(b)(ii)  | <ol style="list-style-type: none"> <li>Three chromosomes;</li> <li>One from each homologous pair;</li> </ol>   | 2     | Ignore shading<br>Only one mark for three chromosomes shown as pairs of chromatids   |

|           |  |   |  |
|-----------|--|---|--|
| 1(b)(iii) | Crossing over / alleles exchanged between chromosomes or chromatids / chiasmata formation / genetic recombination; | 1 | Accept: description of crossing over eg sections of chromatids break and rejoin<br>Neutral: random fertilisation<br>Reject: reference to sister chromatids<br><b>Q</b> Neutral: genes exchanged<br>Neutral: mutation |
|-----------|--|---|--|

| Question | Marking Guidance   | Mark  | Comments   |
|----------|--|-------|--|
| 2(a)     | <ol style="list-style-type: none"> <li>1. Group of similar organisms / organisms with similar features / / organisms with same genes/chromosomes;</li> <li>2. Reproduce / produce offspring;</li> <li>3. That are fertile;</li> </ol>  | 2 max | <ol style="list-style-type: none"> <li>1. Accept: same number of chromosomes</li> <li>1. Accept: smallest taxonomic group</li> <li>1. Reject: genetically identical. Only allow 1 max if mentioned</li> <li>1. <b>Q</b> Neutral: similar genes/chromosomes</li> <li>2. Accept: breed/mate</li> <li>3. Neutral: that are 'viable'</li> </ol> <p>'Produce fertile offspring'<br/>= 2 marks</p> |
| 2(b)(i)  | <p>Correct answer of 6.97 to 7 = 2 marks;</p> <p>One mark for 6320 as numerator or 906 as denominator;</p>   | 2     |  |
| 2(b)(ii) | <ol style="list-style-type: none"> <li>1. Decrease in variety of plants / fewer plant species;</li> <li>2. Fewer habitats/niches;</li> <li>3. Decrease in variety of food / fewer food sources;</li> <li>4. Aspect of clearing forest (killing insects) eg machinery, pesticides;</li> </ol> | 3 max | <ol style="list-style-type: none"> <li>1. Accept: reference to monoculture or description</li> <li>1. Neutral: fewer plants</li> <li>2. Neutral: fewer homes/less shelter</li> <li>3. Neutral: less food</li> <li>3. Accept: less variety of prey</li> <li>4. Neutral: clearing forest unqualified</li> </ol>  |

| Question  | Marking Guidance  | Mark  | Comments  |
|-----------|---|-------|---|
| 3(a)(i)   | <ol style="list-style-type: none"> <li>Groups within groups;</li> <li>No overlap (between groups);</li> </ol>   | 2     | <ol style="list-style-type: none"> <li>Accept: idea of larger groups at the top / smaller groups at the bottom</li> </ol>   |
| 3(a)(ii)  | (Grouped according to) evolutionary links/history/relationships / common ancestry;  | 1     | Neutral: closely related<br>Neutral: genetically similar  |
| 3(b)(i)   | <ol style="list-style-type: none"> <li>(Only) one amino acid different / least differences / similar amino acid sequence / similar primary structure;</li> <li>(So) similar DNA sequence/ base sequence;</li> </ol>                                     | 2     |   |
| 3(b)(ii)  | <ol style="list-style-type: none"> <li>Compared with humans / not compared with each other;</li> <li>Differences may be at different positions / different amino acids affected / does not show where the differences are (in the sequence);</li> </ol> | 1 max | Accept: degenerate code / more than one triplet (codes) for an amino acid   |
| 3(b)(iii) | <ol style="list-style-type: none"> <li>All organisms respire/have cytochrome c;</li> <li>(Cytochrome c structure) is more conserved / less varied (between organisms);</li> </ol>   | 1 max | Accept: converse arguments for haemoglobin <ol style="list-style-type: none"> <li>Accept 'more' instead of 'all'</li> <li>Accept 'animals' instead of organisms'</li> <li>Neutral: cytochrome c is conserved</li> </ol> |

| Question | Marking Guidance   | Mark                          | Comments  |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
|----------|--|-------------------------------|---|--|--|-------------------------------|-------------------------------|-------------------------------|---|---|--|--|---|--|---|--|---|--|---|---|---|-------------------------------|
| 4(a)     | 1. Separates/unwinds/unzips strands/helix / breaks H-bonds;<br>2. (So) <u>nucleotides</u> can attach/are attracted / strands can act as templates;   | 2                             | 1. <b>Q</b> Neutral: strands/helix split<br>1. Accept: unzips bases<br>2. <b>Q</b> Neutral: bases can attach<br>2. Neutral: helix can act as a template   |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
| 4(b)     | <table border="1" data-bbox="316 651 815 1016"> <thead> <tr> <th rowspan="2">Sample</th> <th colspan="3">Type(s) of DNA molecule present in each tube</th> </tr> <tr> <th><math>^{15}\text{N}/^{15}\text{N}</math></th> <th><math>^{15}\text{N}/^{14}\text{N}</math></th> <th><math>^{14}\text{N}/^{14}\text{N}</math></th> </tr> </thead> <tbody> <tr> <td>1</td> <td>✓</td> <td></td> <td></td> </tr> <tr> <td>2</td> <td></td> <td>✓</td> <td></td> </tr> <tr> <td>3</td> <td></td> <td>✓</td> <td>✓</td> </tr> </tbody> </table> | Sample                        | Type(s) of DNA molecule present in each tube  |  |  | $^{15}\text{N}/^{15}\text{N}$ | $^{15}\text{N}/^{14}\text{N}$ | $^{14}\text{N}/^{14}\text{N}$ | 1 | ✓ |  |  | 2 |  | ✓ |  | 3 |  | ✓ | ✓ | 3 | One mark for each correct row |
| Sample   | Type(s) of DNA molecule present in each tube   |                               |   |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
|          | $^{15}\text{N}/^{15}\text{N}$  | $^{15}\text{N}/^{14}\text{N}$ | $^{14}\text{N}/^{14}\text{N}$   |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
| 1        | ✓  |                               |   |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
| 2        |  | ✓                             |   |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
| 3        |  | ✓                             | ✓   |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
| 4(c)(i)  | 1. Similar shape/structure (to cytosine) / added instead of cytosine / binds to guanine;<br>2. Prevents (complementary) base pairing / prevents H-bonds forming / prevents formation of new strand / prevents strand elongation / inhibits/binds to (DNA) polymerase;  | 2                             | 1. Accept: idea that <u>only</u> one group is different<br>1. Reject: same shape<br>2. Accept: prevents cytosine binding<br>Neutral: 'prevents DNA replication' as given in the question stem<br>Neutral: 'competitive inhibitor' unqualified<br>Neutral: inhibits DNA helicase |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |
| 4(c)(ii) | (Cancer cells/DNA) divide/replicate fast(er)/ uncontrollably;  | 1                             | Accept: converse argument for healthy cells   |  |  |                               |                               |                               |   |   |  |  |   |  |   |  |   |  |   |   |   |                               |

| Question | Marking Guidance  | Mark  | Comments  |
|----------|---|-------|---|
| 5(a)(i)  | Prevent cell wall formation / cause (cell) lysis / inhibit ribosomes / inhibit protein synthesis / prevent DNA replication / affect function of cell membrane;  | 1 max | Accept: weaken the cell wall<br>Neutral: damage/break down the cell wall<br><b>Q</b> Reject: if in context of a cellulose cell wall<br>Accept: bind to ribosomes  |
| 5(a)(ii) | (Plasmid/genes transmitted through) cell division/reproduction/replication/generations;   | 1     | Accept: multiply<br>Accept: binary fission<br>Reject: within generations<br>Reject: reference to horizontal gene transmission<br>Reject: mitosis<br>Ignore reference to immunity  |
| 5(b)     | Representative/typical/reliable / different types of bacteria;  | 1     | Neutral: accurate<br>Neutral: reference to anomalies<br><b>Q</b> : Neutral: different strands of bacteria   |
| 5(c)     | (Yes)<br>1. Largest clear zone/diameter/mean (so more bacteria killed);<br>(No)<br>2. Standard deviations of <u>chlorhexidene</u> overlap/share values;<br>3. (Overlap means difference) is not significant / is due to chance; | 3     | Ignore references to methodology<br>2. Neutral: diameters overlap/share values<br>3. Can still be awarded if SD overlap or non-overlap is correctly interpreted<br>3. Accept: (difference) is not real/not reliable<br>3. Neutral: spread is not reliable |
| 5(d)     | 1. <u>Mutation</u> (in bacterium);<br>2. <u>Gene/allele</u> for resistance;   | 2     | 1. Neutral: different strains<br>2. Reject: if in the context of 'immunity'<br>2. Accept: resistant gene/allele   |



| Question                         | Marking Guidance  | Mark      | Comments  |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
|----------------------------------|---|-----------|---|-----------|--------|----------------------------|---|--|--|----------------------------------|---|---|---|-------------------|---|--|--|---|-------------------------------|
| 6(a)                             | <table border="1"> <thead> <tr> <th data-bbox="316 297 523 405">Statement</th> <th data-bbox="523 297 655 405">Haemo-globin</th> <th data-bbox="655 297 807 405">Cellulose</th> <th data-bbox="807 297 919 405">Starch</th> </tr> </thead> <tbody> <tr> <td data-bbox="316 405 523 539">Has a quaternary structure</td> <td data-bbox="523 405 655 539">✓</td> <td data-bbox="655 405 807 539"></td> <td data-bbox="807 405 919 539"></td> </tr> <tr> <td data-bbox="316 539 523 674">Formed by condensation reactions</td> <td data-bbox="523 539 655 674">✓</td> <td data-bbox="655 539 807 674">✓</td> <td data-bbox="807 539 919 674">✓</td> </tr> <tr> <td data-bbox="316 674 523 775">Contains nitrogen</td> <td data-bbox="523 674 655 775">✓</td> <td data-bbox="655 674 807 775"></td> <td data-bbox="807 674 919 775"></td> </tr> </tbody> </table> | Statement | Haemo-globin  | Cellulose | Starch | Has a quaternary structure | ✓ |  |  | Formed by condensation reactions | ✓ | ✓ | ✓ | Contains nitrogen | ✓ |  |  | 3 | One mark for each correct row |
| Statement                        | Haemo-globin  | Cellulose | Starch  |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
| Has a quaternary structure       | ✓   |           |   |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
| Formed by condensation reactions | ✓   | ✓         | ✓   |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
| Contains nitrogen                | ✓   |           |   |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
| 6(b)                             | 16;   | 1         |   |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
| 6(c)                             | <ol style="list-style-type: none"> <li data-bbox="331 898 842 943">1. Higher affinity / loads more oxygen;</li> <li data-bbox="331 943 882 987">2. At low/same/high partial pressure/pO<sub>2</sub>;</li> <li data-bbox="331 987 858 1032">3. Oxygen moves from mother/to fetus;</li> </ol>   | 2 max     |   |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
| 6(d)                             | <ol style="list-style-type: none"> <li data-bbox="331 1077 810 1122">1. Low affinity / oxygen dissociates;</li> <li data-bbox="331 1122 667 1200">2. (Oxygen) to respiring tissues/muscles/cells;</li> </ol>  | 2         | Assume 'it' is adult haemoglobin <ol style="list-style-type: none"> <li data-bbox="1118 1167 1417 1267">1. Accept: converse if 'fetal haemoglobin' is clearly stated</li> <li data-bbox="1118 1279 1305 1346">2. <b>Q:</b> Neutral 'respire'</li> </ol> |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |
| 6(e)                             | Enough adult Hb produced / enough oxygen released / idea that curves/affinities/Hb are similar / more red blood cells produced;   | 1         | Neutral: 'adult Hb is also produced' as in the question stem<br><br>Reject: curves/affinities/Hb are the same   |           |        |                            |   |  |  |                                  |   |   |   |                   |   |  |  |   |                               |

| Question | Marking Guidance  | Mark  | Comments   |
|----------|---|-------|--|
| 7(a)     | <ol style="list-style-type: none"> <li>1. Population formed by a small number of founders/people /30 people;</li> <li>2. (Founders show) less genetic diversity / small(er) gene pool / less variety of alleles;</li> <li>3. Individuals breed within group / do not breed with outsiders;</li> <li>4. High(er) chance of inheriting <u>allele</u> (than in non-Amish population);</li> </ol> | 3 max | <p>Accept: converse arguments for the non-Amish population</p> <ol style="list-style-type: none"> <li>2. <b>Q</b> Neutral: fewer alleles</li> <li>3. Accept: inbreeding for 'individuals breed within group'</li> <li>3. Accept: do not interbreed</li> <li>3. <b>Q</b> Reject: interbreeding for 'individuals breed within group'</li> <li>4. Do not award for 'allele passed on' only</li> </ol> |
| 7(b)     | 250 000;  | 1     |  |
| 7(c)(i)  | <p>Loss of 3 bases/triplet = 2 marks;;</p> <p>Loss of base(s) = 1 mark;</p>   | 2     | <p>'Stop codon/code formed' = 1 mark max unless related to the last amino acid</p> <p>eg triplet for last amino acid is changed to a stop codon/code = 2 marks</p> <p>3 bases/triplet forms an intron = 2 marks</p> <p>Accept: descriptions for 'intron' eg non-coding DNA</p> <p>'Loss of codon' = 2 marks</p>  |
| 7(c)(ii) | <ol style="list-style-type: none"> <li>1. Change in tertiary structure/ active site;</li> <li>2. (So) faulty/non-functional protein /enzyme;</li> </ol>   | 2     | <p>Neutral: change in 3D shape/ structure</p> <p>Accept: reference to examples of loss of function eg fewer E-S complexes formed</p>   |

| Question | Marking Guidance  | Mark  | Comments   |
|----------|---|-------|--|
| 8(a)     | <p>(In the root)</p> <ol style="list-style-type: none"> <li>Casparian strip blocks apoplast pathway / only allows symplast pathway;</li> <li>Active transport by <u>endodermis</u>;</li> <li>(Of) ions/salts into xylem;</li> <li>Lower water potential in xylem / water enters xylem by osmosis /down a water potential gradient;</li> </ol> <p>(Xylem to leaf)</p> <ol style="list-style-type: none"> <li>Evaporation / transpiration (from leaves);</li> <li>(Creates) cohesion / tension / H-bonding between water molecules / negative pressure;</li> <li>Adhesion / water molecules bind to xylem;</li> <li>(Creates continuous) water column;</li> </ol> | 6 max | <p>Assume all points are in the correct location unless context suggests otherwise</p> <p>4. <b>Q</b> Neutral: 'along' a water potential gradient</p> <p>'Transpiration pull' = 2 marks (5. &amp; 6.)</p> <p>6. Accept 'pulling'</p> <p>6. <b>Q</b> Neutral: 'suction'</p> |
| 8(b)     | <p>Correct answer of 342.8-343 = 2 marks;;</p> <p>Credit incorrect answers that show the numerator as 144 (or 186-42) or denominator as 42 for 1 mark;</p>  | 2     |  |
| 8(c)     | <ol style="list-style-type: none"> <li>More air/oxygen enters / air/oxygen enters quickly/quicker;</li> <li>(So) maintains/greater diffusion or concentration <u>gradient</u>;</li> </ol>   | 2     | <ol style="list-style-type: none"> <li>Accept: converse for carbon dioxide</li> <li>Can be in any correct context eg insect, tracheoles, muscle</li> <li>Neutral: air/oxygen enters</li> </ol>   |
| 8(d)     | <p>Large(r) SA:VOL / short(er) <u>diffusion</u> distance (to tissues);</p>  | 1     | Accept: thin diffusion pathway   |
| 8(e)     | <p>6 / 6.6 / 6.7 / 7 / 7.5 / 8 = 2 marks;;</p> <p>Award 1 mark for incorrect answers that have divided 60 by any number;</p>  | 2     | Different answers given for different interpretations of the graph   |

|      |   |       |  |
|------|---|-------|--|
| 8(f) | Less/no water lost / (more) water retained;   | 1     | <p>Accept: less dehydration / less evaporation</p> <p><b>Q</b> Reject: less 'transpiration'</p> <p><b>Q</b> Reject: less water lost by osmosis</p>   |
| 8(g) | <ol style="list-style-type: none"> <li>1. Greater <u>surface area</u> exposed to air;</li> <li>2. Gases move/diffuse faster in air than through water;</li> <li>3. Increases volume/amount of air;</li> </ol> | 1 max | <p>Neutral: shorter diffusion distance</p> <ol style="list-style-type: none"> <li>2. <b>Q</b> Neutral: 'harder to diffuse'</li> <li>2. Accept gases diffuse directly, rather than through water</li> </ol> |

| Question | Marking Guidance   | Mark  | Comments   |
|----------|--|-------|--|
| 9(a)     | <p>Any two suitable suggestions eg</p> <ol style="list-style-type: none"> <li>1. Volume/concentration of skin lipid;</li> <li>2. Age/sexual maturity;</li> <li>3. <u>Species</u> of snake;</li> <li>4. Size of <u>male</u>;</li> <li>5. Colour;</li> <li>6. Temperature;</li> <li>7. Light;</li> <li>8. Time of day/year/breeding season;</li> <li>9. Duration/length of time observing;</li> <li>10. Diet;</li> <li>11. Filter paper;</li> <li>12. Size of cage;</li> </ol> | 2 max | <p>1. Accept: amount</p> <p>Neutral: environment / health / body mass / number of snakes</p>   |
| 9(b)     | To avoid bias;   | 1     |  |
| 9(c)     | <ol style="list-style-type: none"> <li>1. To avoid change in (courtship) behaviour (due to past experience);</li> <li>2. To observe a typical/general/representative (response);</li> </ol>  | 1 max | <p>Accept: ethical arguments eg causing distress to snakes</p> <p>Neutral: reference to anomalous results</p>                            |
| 9(d)     | Filter paper without (skin) lipids / untreated filter paper / filter paper with water / (female) snakes with lipids removed;   | 1     | <p>Accept: filter paper qualified eg <i>only</i> filter paper</p> <p>Neutral: reference to using male snakes/lipids from male snakes</p> |
| 9(e)     | <ol style="list-style-type: none"> <li>1. Similar response to lipids and (whole) snakes;</li> <li>2. (So males are) responding to lipids;</li> <li>3. (So males are) not responding to (whole) snakes/visual clues;</li> </ol>   | 2 max | Neutral: greater response to long snakes and lipids from long snakes as in the question stem   |

|      |   |       |  |
|------|---|-------|--|
| 9(f) | <p>(Parent/offspring)</p> <ol style="list-style-type: none"> <li>1. Produce more/larger offspring/eggs;</li> <li>2. Better predators / fitter / more successful at gaining food / less likely to be eaten / more able to protect offspring/eggs;</li> <li>3. (More) sexually mature / fertile;</li> <li>4. Have more food stores for offspring/eggs;</li> </ol> | 2 max | 3. Neutral: mature   |
| 9(g) | <ol style="list-style-type: none"> <li>1. (Males) respond to/sense (unsaturated) <u>fatty acids</u>;</li> <li>2. (Long females) produce/have more fatty acids / positive correlation;</li> </ol>  | 2     | <ol style="list-style-type: none"> <li>1. Reference to courtship behaviour on its own is not sufficient</li> </ol> <p>Reference to 'lipids/fats' is neutral for both mark points. However, if fatty acids are mentioned for either mark point, accept lipids/fats = fatty acids for the other mark point</p> |
| 9(h) | <ol style="list-style-type: none"> <li>1. Draw a line of best fit;</li> <li>2. Extrapolation / extend line;</li> </ol>  | 2     |  |
| 9(i) | Results vary for a particular body size/% / values overlap / small sample size / idea of reaching maximum/100%/ a plateau;  | 1     | <p>Neutral: reference to inaccurate line of best fit</p> <p>Neutral: 'results vary'</p>  |
| 9(j) | (Other females/species) produce different fatty acids;  | 1     | <p>Must refer to fatty acids rather than just 'lipids/fats'</p> <p>Accept: lack of receptors</p>   |