



Higher Biology:

Course content change comparison

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Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
<p>1 The structure of DNA</p> <p>(a) Structure of DNA — nucleotides (deoxyribose sugar, phosphate and base), sugar–phosphate backbone, base pairing (adenine–thymine and guanine–cytosine), by hydrogen bonds and double stranded antiparallel structure, with deoxyribose and phosphate at 3' and 5' ends of each strand respectively, forming a double helix.</p>	<p>The base sequence of DNA forms the genetic code.</p>	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
<p>(b) Organisation of DNA — Prokaryotes have a single, circular chromosome and smaller circular plasmids.</p> <p>Eukaryotes all have linear chromosomes, in the nucleus, which are tightly coiled and packaged with associated proteins. They also contain circular chromosomes in their mitochondria and chloroplasts. Yeast is a special example of a eukaryote as it also has plasmids.</p>	<p>The associated proteins are called histones.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>

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DNA and the genome		
<p>2 Replication of DNA (a) Replication of DNA by DNA polymerase and primers.</p> <p>DNA polymerase adds DNA nucleotides, using complementary base pairing, to the deoxyribose (3') end of the new DNA strand which is forming.</p> <p>Fragments of DNA are joined together by ligase.</p>	<p>Prior to cell division, DNA is replicated by a DNA polymerase. DNA polymerase needs primers to start replication. A primer is a short strand of nucleotides which binds to the 3' end of the template DNA strand allowing polymerase to add DNA nucleotides.</p> <p>DNA is unwound and hydrogen bonds between bases are broken to form two template strands. DNA polymerase can only add DNA nucleotides in one direction resulting in the leading strand being replicated continuously and the lagging strand replicated in fragments.</p>	<p>No change to content.</p> <p>Minor rewording for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>(b) Polymerase chain reaction (PCR) amplifies DNA using complementary primers for specific target sequences.</p>	<p>In PCR, primers are short strands of nucleotides which are complementary to specific target sequences at the two ends of the region of DNA to be amplified.</p>	<p>Minor rewording for clarification.</p> <p>Clarification of required depth of knowledge.</p>

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<p>Repeated cycles of heating and cooling amplify the target region of DNA.</p> <p>Practical applications of PCR.</p>	<p>DNA is heated to between 92 and 98 °C to separate the strands. It is then cooled to between 50 and 65 °C to allow primers to bind to target sequences. It is then heated to between 70 and 80 °C for heat-tolerant DNA polymerase to replicate the region of DNA.</p> <p>PCR can amplify DNA to help solve crimes, settle paternity suits and diagnose genetic disorders.</p>	<p>Clarification of required depth of knowledge.</p> <p>Positive and negative controls removed.</p>
<p>3 Gene expression (a) Gene expression involves the transcription and translation of DNA sequences.</p> <p>Transcription and translation involves three types of RNA (mRNA, tRNA and rRNA).</p> <p>Messenger RNA (mRNA) carries a copy of the DNA code from the nucleus to the ribosome.</p>	<p>Only a fraction of the genes in a cell are expressed.</p> <p>RNA is single stranded and is composed of nucleotides containing ribose sugar, phosphate and one of four bases: cytosine, guanine, adenine and uracil.</p> <p>mRNA is transcribed from DNA in the nucleus and translated into proteins by ribosomes in the cytoplasm. Each triplet of bases on the mRNA molecule is called a codon and codes for a specific amino acid.</p>	<p>New key area heading.</p> <p>Former key areas 1.3 (a) and (b) combined.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>Information on phenotype determination moved to key area 1.3 (e).</p> <p>Influence by intra- and extra-cellular environmental factors removed.</p>

Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
<p>Transfer RNA (tRNA) folds due to complementary base pairing. Each tRNA molecule carries its specific amino acid to the ribosome.</p> <p>Ribosomal RNA (rRNA) and proteins form the ribosome.</p>	<p>A tRNA molecule has an anticodon (an exposed triplet of bases) at one end and an attachment site for a specific amino acid at the other end.</p>	
<p>(b) The role of RNA polymerase in transcription of DNA into primary mRNA transcripts.</p> <p>RNA splicing forms a mature mRNA transcript.</p> <p>The introns of the primary transcript are non-coding regions and are removed.</p> <p>The exons are coding regions and are joined together to form the mature transcript.</p>	<p>RNA polymerase moves along DNA unwinding the double helix and breaking the hydrogen bonds between the bases. RNA polymerase synthesises a primary transcript of mRNA from RNA nucleotides by complementary base pairing.</p> <p>Uracil in RNA is complementary to adenine.</p> <p>The order of the exons is unchanged during splicing.</p>	<p>Formerly key area 1.3 (c).</p> <p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>

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DNA and the genome		
<p>(c) tRNA is involved in the translation of mRNA into a polypeptide at a ribosome. Translation begins at a start codon and ends at a stop codon. Anticodons bond to codons by complementary base pairing, translating the genetic code into a sequence of amino acids. Peptide bonds join the amino acids together. Each tRNA then leaves the ribosome as the polypeptide is formed.</p>		<p>Formerly key area 1.3 (d).</p> <p>Structure of tRNA moved to key area 1.3 (a).</p> <p>Reworded for clarification.</p>
<p>(d) Different proteins can be expressed from one gene, as a result of alternative RNA splicing. Different mature mRNA transcripts are produced from the same primary transcript depending on which exons are retained.</p>		<p>Formerly key area 1.3 (e).</p> <p>Reworded for clarification.</p> <p>Post-translational modification removed.</p>
<p>(e) Amino acids are linked by peptide bonds to form polypeptides. Polypeptide chains fold to form the three-dimensional shape of a protein, held together by hydrogen bonds and other interactions between individual amino acids. Proteins have a large variety of shapes which determines their functions.</p> <p>Phenotype is determined by the proteins produced as the result of gene expression.</p>	<p>Details of other interactions and levels of protein structure are not required.</p> <p>Environmental factors also influence phenotype.</p>	<p>Formerly key area 1.3 (f).</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>Information on phenotype from key area 1.3 (a) included here.</p>

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DNA and the genome		
<p>4 Cellular differentiation</p> <p>(a) Cellular differentiation is the process by which a cell expresses certain genes to produce proteins characteristic for that type of cell. This allows a cell to carry out specialised functions.</p> <p>Differentiation into specialised cells from meristems in plants; embryonic and tissue (adult) stem cells in animals.</p>	<p>Meristems are regions of unspecialised cells in plants that can divide (self-renew) and/or differentiate.</p> <p>Stem cells are unspecialised cells in animals that can divide (self-renew) and/or differentiate.</p> <p>There is no requirement to learn examples of differentiated animal and plant cells.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>(b) Embryonic and tissue stem cells.</p> <p>Cells in the very early embryo can differentiate into all the cell types that make up the organism and so are pluripotent.</p> <p>Tissue stem cells are involved in the growth, repair and renewal of the cells found in that tissue. They are multipotent.</p>	<p>All the genes in embryonic stem cells can be switched on so these cells can differentiate into any type of cell.</p> <p>Tissue stem cells are multipotent as they can differentiate into all of the types of cell found in a particular tissue type. For example, blood stem cells located in bone marrow can give rise to all types of blood cell.</p>	<p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
<p>DNA and the genome</p> <p>Therapeutic and research uses of stem cells.</p> <p>Therapeutic uses involve the repair of damaged or diseased organs or tissues.</p> <p>Research uses involve stem cells being used as model cells to study how diseases develop or being used for drug testing.</p> <p>The ethical issues of using embryonic stem cells.</p>	<p>The therapeutic uses of stem cells should be exemplified by how they are used in corneal repair and the regeneration of damaged skin.</p> <p>Stem cells from the embryo can self-renew, under the right conditions, in the lab.</p> <p>Stem cell research provides information on how cell processes such as cell growth, differentiation and gene regulation work.</p> <p>Use of embryonic stem cells can offer effective treatments for disease and injury; however, it involves destruction of embryos.</p>	<p>Regulation of stem cell use removed.</p>
<p>5 The structure of the genome</p> <p>The genome of an organism is its entire hereditary information encoded in DNA.</p> <p>A genome is made up of genes and other DNA sequences that do not code for proteins.</p> <p>DNA sequences that code for protein are defined as genes. Other sequences regulate transcription and others are transcribed but never translated.</p>	<p>Most of the eukaryotic genome consists of non-coding sequences.</p> <p>Details of regulation of transcription (eg Jacob–Monod hypothesis) not required.</p> <p>tRNA and rRNA are non-translated forms of RNA.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
<p>6 Mutations</p> <p>(a) Mutations are changes in the DNA that can result in no protein or an altered protein being synthesised.</p>		<p>No change to content.</p> <p>Minor rewording for clarification.</p>
<p>(b) Single gene mutations involve the alteration of a DNA nucleotide sequence as a result of the substitution, insertion or deletion of nucleotides.</p> <p>Nucleotide substitutions — missense, nonsense and splice-site mutations.</p> <p>Nucleotide insertions or deletions result in frame-shift mutations.</p>	<p>Missense mutations result in one amino acid being changed for another. This may result in a non-functional protein or have little effect on the protein.</p> <p>Nonsense mutations result in a premature stop codon being produced which results in a shorter protein.</p> <p>Splice-site mutations result in some introns being retained and/or some exons not being included in the mature transcript.</p> <p>Frame-shift mutations cause all of the codons and all of the amino acids after the mutation to be changed. This has a major effect on the structure of the protein produced.</p>	<p>Clarification of required depth of knowledge.</p> <p>Expansion of nucleotide sequence repeat removed.</p>

Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
(c) Chromosome structure mutations — duplication, deletion, inversion and translocation.	<p>Duplication is where a section of a chromosome is added from its homologous partner.</p> <p>Deletion is where a section of a chromosome is removed.</p> <p>Inversion is where a section of chromosome is reversed.</p> <p>Translocation is where a section of a chromosome is added to a chromosome, not its homologous partner.</p> <p>The substantial changes in chromosome mutations often make them lethal.</p>	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
(d) Importance of mutations and gene duplication in evolution.	<p>Duplication allows potential beneficial mutations to occur in a duplicated gene whilst the original gene can still be expressed to produce its protein.</p>	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
		Former key area 1.6 (e) removed.

Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
<p>7 Evolution (a) Evolution — the changes in organisms over generations as a result of genomic variations.</p>		<p>No change to content.</p>
<p>(b) Selection. Natural selection is the non-random increase in frequency of DNA sequences that increase survival and the non-random reduction in the frequency of deleterious sequences.</p> <p>The changes in phenotype frequency as a result of stabilising, directional and disruptive selection.</p> <p>Natural selection is more rapid in prokaryotes. Prokaryotes can exchange genetic material horizontally, resulting in faster evolutionary change than in organisms that only use vertical transfer.</p>	<p>In stabilising selection, an average phenotype is selected for and extremes of the phenotype range are selected against. In directional selection, one extreme of the phenotype range is selected for. In disruptive selection, two or more phenotypes are selected for.</p> <p>Horizontal gene transfer is where genes are transferred between individuals in the same generation. Methods of horizontal transfer are not required.</p>	<p>Former key areas 1.7 (b) and (c) combined.</p> <p>Reworded for clarification.</p> <p>Sexual selection removed.</p> <p>Clarification of required depth of knowledge.</p> <p>Transfer of DNA sequences between prokaryotes and eukaryotes removed.</p>

Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
	Vertical gene transfer is where genes are transferred from parent to offspring as a result of sexual or asexual reproduction.	
(c) Speciation. Speciation is the generation of new biological species by evolution as a result of isolation, mutation and selection. The importance of isolation barriers in preventing gene flow between sub-populations during speciation. Geographical barriers lead to allopatric speciation and behavioural or ecological barriers lead to sympatric speciation.	A species is a group of organisms capable of interbreeding and producing fertile offspring, and which does not normally breed with other groups.	Formerly key area 1.7 (e). Minor rewording for clarification. Clarification of required depth of knowledge. Hybrid zones removed.
		Former key area 1.7 (d) removed.
8 Genomic sequencing (a) In genomic sequencing the sequence of nucleotide bases can be determined for individual genes and entire genomes. Comparison of genomes from different species. Comparison of genomes reveals that many genes are highly conserved across different organisms.	Computer programs can be used to identify base sequences by looking for sequences similar to known genes. To compare sequence data, computer and statistical analyses (bioinformatics) are required. Many genomes have been sequenced, particularly of disease-causing organisms, pest species and species that are important model organisms for research.	Former key areas 1.8 (a) and (c) combined. No change to content. Clarification of required depth of knowledge.

Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
<p>(b) Evidence from phylogenetics and molecular clocks to determine the main sequence of events in evolution. The sequence of events can be determined using sequence data and fossil evidence.</p> <p>Comparison of sequences provides evidence of the three domains of life — bacteria, archaea and eukaryotes.</p>	<p>Phylogenetics is the study of evolutionary history and relationships.</p> <p>Use of sequence data to study the evolutionary relatedness among groups of organisms. Sequence divergence is used to estimate time since lineages diverged.</p> <p>Use of sequence data and fossil evidence to determine the main sequence of events in evolution of life: cells, last universal ancestor, prokaryotes, photosynthetic organisms, eukaryotes, multicellularity, animals, vertebrates, land plants.</p> <p>Molecular clocks are used to show when species diverged during evolution. They assume a constant mutation rate and show differences in DNA sequences or amino acid sequences. Therefore differences in sequence data between species indicate the time of divergence from a common ancestor.</p>	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
<p>(c) An individual's genome can be analysed to predict the likelihood of developing certain diseases.</p> <p>Pharmacogenetics and personalised medicine.</p>	<p>Pharmacogenetics is the use of genome information in the choice of drugs.</p>	<p>Formerly key area 1.8 (d).</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
DNA and the genome		
	An individual's personal genome sequence can be used to select the most effective drugs and dosage to treat their disease (personalised medicine).	Difficulties with personalised medicine removed.

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
<p>1 Metabolic pathways</p> <p>(a) Metabolic pathways are integrated and controlled pathways of enzyme-catalysed reactions within a cell.</p> <p>Metabolic pathways can have reversible steps, irreversible steps and alternative routes.</p> <p>Reactions within metabolic pathways can be anabolic or catabolic. Anabolic reactions build up large molecules from small molecules and require energy. Catabolic reactions break down large molecules into smaller molecules and release energy.</p>		<p>New key area heading.</p> <p>Former key areas 2.1 (a) and 2.1 (a) (i) combined.</p> <p>No change to content.</p> <p>Minor rewording for clarification.</p>
<p>(b) Protein pores, pumps and enzymes are embedded in membranes.</p>	<p>No requirement to know details of sodium potassium pump.</p>	<p>Formerly part of key area 2.1 (a) (ii).</p> <p>Section on membrane compartments removed.</p>
<p>(c) Metabolic pathways are controlled by the presence or absence of particular enzymes and the regulation of the rate of reaction of key enzymes.</p> <p>Induced fit and the role of the active site of an enzyme in affecting activation energy and the affinity of the substrate and products for the active site.</p>	<p>Induced fit occurs when the active site changes shape to better fit the substrate after the substrate binds.</p>	<p>Former key areas 2.1 (b) (i) and (ii) combined.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
<p>The effects of substrate and product concentration on the direction and rate of enzyme reactions.</p> <p>Control of metabolic pathways through competitive, non-competitive and feedback inhibition of enzymes.</p>	<p>The substrate molecule(s) have a high affinity for the active site and the subsequent products have a low affinity, allowing them to leave the active site. Some metabolic reactions are reversible and the presence of a substrate or the removal of a product will drive a sequence of reactions in a particular direction.</p> <p>Competitive inhibitors bind at the active site preventing the substrate from binding. Competitive inhibition can be reversed by increasing substrate concentration.</p> <p>Non-competitive inhibitors bind away from the active site but change the shape of the active site preventing the substrate from binding. Non-competitive inhibition cannot be reversed by increasing substrate concentration.</p> <p>Feedback inhibition occurs when the end-product in the metabolic pathway reaches a critical concentration. The end-product then inhibits an earlier enzyme, blocking the pathway, and so prevents further synthesis of the end-product.</p>	<p>Multi-enzyme complexes removed.</p>

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
<p>2 Cellular respiration (a) Metabolic pathways of cellular respiration.</p> <p>Glycolysis is the breakdown of glucose to pyruvate in the cytoplasm.</p> <p>ATP is required for the phosphorylation of glucose and intermediates during the energy investment phase of glycolysis. This leads to the generation of more ATP during the energy pay-off stage and results in a net gain of ATP.</p> <p>In aerobic conditions, pyruvate is broken down to an acetyl group that combines with coenzyme A forming acetyl coenzyme A.</p> <p>In the citric acid cycle the acetyl group from acetyl coenzyme A combines with oxaloacetate to form citrate. During a series of enzyme controlled steps, citrate is gradually converted back into oxaloacetate which results in the generation of ATP and release of carbon dioxide.</p> <p>The citric acid cycle occurs in the matrix of the mitochondria.</p> <p>Dehydrogenase enzymes remove hydrogen ions and electrons and pass them to the coenzyme NAD, forming NADH. This occurs in both glycolysis and the citric acid cycle.</p>		<p>New key area heading.</p> <p>Former key areas 2.2 (a) and (c) combined.</p> <p>Reworded for clarification.</p> <p>Section on pathway when oxygen absent moved to key area 2.2 (c).</p> <p>FAD removed.</p> <p>Electrons being referred to as 'high energy' removed.</p>

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
The hydrogen ions and electrons from NADH are passed to the electron transport chain on the inner mitochondrial membrane.		
<p>(b) ATP synthesis — electrons are passed along the electron transport chain releasing energy.</p> <p>This energy allows hydrogen ions to be pumped across the inner mitochondrial membrane. The flow of these ions back through the membrane protein ATP synthase results in the production of ATP.</p> <p>Finally, hydrogen ions and electrons combine with oxygen to form water.</p>	<p>The electron transport chain is a series of carrier proteins attached to the inner mitochondrial membrane.</p>	<p>Former key area 2.2 (d).</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>Reference to electrons being 'high energy' removed.</p>
<p>(c) Fermentation.</p> <p>In the absence of oxygen, fermentation takes place in the cytoplasm.</p> <p>In animal cells, pyruvate is converted to lactate in a reversible reaction.</p> <p>In plants and yeast, ethanol and CO₂ are produced in an irreversible reaction.</p> <p>Fermentation results in much less ATP being produced than in aerobic respiration.</p>		<p>Formerly part of key area 2.2 (c).</p> <p>Reworded for clarification.</p>
(d) The role of ATP in the transfer of energy.	ATP is used to transfer energy to cellular processes which require energy.	Formerly part of key area 2.2 (b). Clarification of required depth of knowledge.

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
		Former key area 2.2 (e) removed.
<p>3 Metabolic rate</p> <p>(a) Measurement of oxygen consumption, carbon dioxide and heat production to compare metabolic rates.</p>	Metabolic rate can be measured using respirometers, oxygen probes, carbon dioxide probes and calorimeters.	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
<p>(b) Organisms with high metabolic rates require more efficient delivery of oxygen to cells.</p> <p>Comparative anatomy and physiology of heart chambers and circulatory systems in amphibians, reptiles, mammals and birds, and heart and circulation in fish.</p>	<p>Birds and mammals have higher metabolic rates than reptiles and amphibians, which in turn have higher metabolic rates than fish.</p> <p>Birds and mammals have a complete double circulatory system consisting of two atria and two ventricles. Amphibians and most reptiles have an incomplete double circulatory system consisting of two atria and one ventricle. Fish have a single circulatory system consisting of one atrium and one ventricle.</p> <p>Complete double circulatory systems enable higher metabolic rates to be maintained. There is no mixing of oxygenated and deoxygenated blood and the oxygenated blood can be pumped out at a higher pressure. This enables more efficient oxygen delivery to cells.</p>	<p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>'Lung arrangement' removed.</p>

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
		Former key areas 2.3 (c) and 2.3 (d) removed.
<p>4 Metabolism in conformers and regulators</p> <p>(a) The ability of an organism to maintain its metabolic rate is affected by external abiotic factors.</p>	Abiotic factors — temperature, salinity and pH.	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
<p>(b) Conformers' internal environment is dependent upon external environment. Conformers use behavioural responses to maintain optimum metabolic rate. Conformers have low metabolic costs and a narrow range of ecological niches.</p>	Behavioural responses by conformers allow them to tolerate variation in their external environment to maintain optimum metabolic rate.	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>(c) Regulators maintain their internal environment regardless of external environment.</p> <p>Regulators use metabolism to control their internal environment, which increases the range of possible ecological niches.</p> <p>This regulation requires energy to achieve homeostasis. This increases their metabolic costs.</p>		<p>No change to content.</p> <p>Reworded for clarification.</p>

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
<p>(d) Thermoregulation by negative feedback — the role of the hypothalamus, nerves and effectors.</p> <p>The role of corrective responses to an increase in body temperature — sweating, vasodilation of blood vessels and decreased metabolic rate.</p> <p>The corrective responses to a decrease in body temperature — shivering, vasoconstriction of blood vessels, hair erector muscles contracting and increased metabolic rate.</p>	<p>The hypothalamus is the temperature monitoring centre. Information is communicated by electrical impulses through nerves to the effectors, which bring about corrective responses to return temperature to normal.</p> <p>Sweating — body heat used to evaporate water in the sweat, cooling the skin. Vasodilation — increased blood flow to the skin increases heat loss. Decreased metabolic rate — less heat produced.</p> <p>Shivering — muscle contraction generates heat. Vasoconstriction — decreased blood flow to skin decreases heat loss. Hair erector muscles contract — traps layer of insulating air. Increased metabolic rate — more heat produced.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>(e) Importance of regulating temperature (thermoregulation) for optimal enzyme activity and high diffusion rates to maintain metabolism.</p>		<p>No change to content.</p> <p>Reworded for clarification.</p>

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Metabolism and survival		
<p>5 Metabolism and adverse conditions Many environments vary beyond the tolerable limits for normal metabolic activity for any particular organism. Some animals have adapted to survive these adverse conditions while others avoid them.</p> <p>(a) Surviving adverse conditions by dormancy.</p> <p>Dormancy is part of some organisms' lifecycle to allow survival during a period when the costs of continued normal metabolic activity would be too high. The metabolic rate can be reduced during dormancy to save energy.</p> <p>Dormancy can be predictive or consequential.</p> <p>Some mammals survive during winter/low temperatures by hibernating. Aestivation allows survival in periods of high temperature or drought. Daily torpor is a period of reduced activity in some animals with high metabolic rates.</p>	<p>During dormancy there is a decrease in metabolic rate, heart rate, breathing rate and body temperature.</p> <p>Predictive dormancy occurs before the onset of adverse conditions. Consequential dormancy occurs after the onset of adverse conditions.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>(b) Avoiding adverse conditions by migration.</p> <p>Migration avoids metabolic adversity by expending energy to relocate to a more suitable environment.</p>		<p>No change to content.</p> <p>Reworded for clarification.</p>

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Metabolism and survival		
<p>Migratory behaviour can be innate and learned.</p> <p>Specialised techniques are used to study long-distance migration.</p>	<p>Examples of specialist techniques are: satellite tracking and leg rings.</p>	<p>Clarification of required depth of knowledge.</p>
		<p>Former key area 2.5 (c) removed.</p>
<p>6 Environmental control of metabolism</p> <p>Micro-organisms are archaea, bacteria and some species of eukaryotes.</p>	<p>Micro-organisms use a wide variety of substrates for metabolism and produce a range of products from their metabolic pathways.</p> <p>Micro-organisms are used because of their adaptability, ease of cultivation and speed of growth.</p>	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
<p>(a) Variations in growth media and control of environmental factors.</p> <p>When culturing micro-organisms, their growth media requires raw materials for biosynthesis as well as an energy source.</p>	<p>Many micro-organisms produce all the complex molecules required for biosynthesis, for example amino acids, vitamins and fatty acids. Other micro-organisms require these to be supplied in the growth media.</p> <p>An energy source is derived either from chemical substrates or from light in photosynthetic micro-organisms.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge</p>

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Metabolism and survival		
<p>Culture conditions: Sterility Control of temperature, oxygen levels and pH.</p>	<p>Sterile conditions in fermenters reduce competition with desired micro-organisms for nutrients and reduce the risk of spoilage of the product.</p>	
<p>(b) Phases of growth and changes in culture conditions. Phases — lag, log/exponential, stationary and death.</p>	<p>The lag phase is where enzymes are induced to metabolise substrates.</p> <p>The log/exponential phase contains the most rapid growth of micro-organisms due to plentiful nutrients.</p> <p>The stationary phase occurs due to the nutrients in the culture media becoming depleted and the production of toxic metabolites. Secondary metabolites are also produced, such as antibiotics. In the wild these metabolites confer an ecological advantage by allowing the micro-organisms which produce them to outcompete other micro-organisms.</p> <p>The death phase occurs due to the toxic accumulation of metabolites or the lack of nutrients in the culture.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
Growth curves of micro-organisms. Viable and total cell count.	Use of semi-logarithmic scales in producing or interpreting growth curves of micro-organisms. Viable cell counts involve counting only the living micro-organisms whereas total cell counts involve counting viable and dead cells. Only viable cell counts show a death phase where cell numbers are decreasing.	
		Former key area 2.6 (c) removed.
7 Genetic control of metabolism (a) Wild strains of micro-organisms can be improved by mutagenesis, or recombinant DNA technology.	Exposure to UV light and other forms of radiation or mutagenic chemicals results in mutations, some of which may produce an improved strain of micro-organism.	Selective breeding and culture removed. Bacteria transferring DNA removed. Sexual reproduction in fungi removed. Reworded for clarification. Clarification of required depth of knowledge.
(b) Recombinant DNA technology involves the use of recombinant plasmids and artificial chromosomes as vectors.	A vector is a DNA molecule used to carry foreign genetic information into another cell and both plasmids and artificial chromosomes are used as vectors during recombinant DNA technology.	Reworded for clarification. Clarification of required depth of knowledge.

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
<p>The role of the enzymes restriction endonucleases and ligase in recombinant DNA technology.</p> <p>Recombinant plasmids and artificial chromosomes contain restriction sites, regulatory sequences, an origin of replication and selectable markers.</p>	<p>Artificial chromosomes are preferable to plasmids as vectors when larger fragments of foreign DNA are required to be inserted.</p> <p>Restriction endonucleases cut open plasmids and specific genes out of chromosomes, leaving sticky ends.</p> <p>Complementary sticky ends are produced when the same restriction endonuclease is used to cut open the plasmid and the gene from the chromosome. Ligase seals the gene into the plasmid.</p> <p>Restriction sites contain target sequences of DNA where specific restriction endonucleases cut.</p> <p>Regulatory sequences control gene expression and origin of replication allows self-replication of the plasmid/artificial chromosome.</p> <p>Selectable markers such as antibiotic resistance genes protect the micro-organism from a selective agent (antibiotic) that would normally kill it or prevent it growing.</p> <p>Selectable marker genes present in the vector ensure that only micro-organisms that have taken up the vector grow in the presence of the selective agent (antibiotic).</p>	<p>Genes can be introduced to remove inhibitory controls or amplify metabolic steps removed.</p> <p>Control of gene expression in recombinant plasmids and artificial chromosomes removed.</p>

Key areas	Depth of knowledge required	Changes to content
Metabolism and survival		
<p>As a safety mechanism, genes are often introduced that prevent the survival of the micro-organism in an external environment.</p> <p>Use of recombinant yeast cells to produce active forms of the protein which are inactive in bacteria.</p>	<p>Recombinant yeast cells may be used as plant or animal recombinant DNA expressed in bacteria may result in polypeptides being incorrectly folded.</p>	<p>Reference to post-translational modifications removed.</p> <p>Former key area 2.8 removed. This will become part of the Suggested Learning Activities for key area 2.7.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>1 Food supply, plant growth and productivity</p> <p>(a) Food supply. Food security and sustainable food production. Increase in human population and concern for food security leads to a demand for increased food production. Food production must be sustainable and not degrade the natural resources on which agriculture depends.</p> <p>Agricultural production depends on factors that control photosynthesis and plant growth. The area to grow crops is limited. Increased food production will depend on factors that control plant growth — breeding of higher yielding cultivars, use of fertiliser, protecting crops from pests, diseases and competition.</p> <p>Livestock produce less food per unit area than crop plants due to loss of energy between trophic levels. Livestock production is often possible in habitats unsuitable for growing crops.</p>	<p>Food security is the ability of human populations to access food of sufficient quality and quantity.</p> <p>All food production is dependent ultimately upon photosynthesis. Plant crop examples include cereals, potato, roots and legumes. Breeders seek to develop crops with higher nutritional values, resistance to pests and diseases, physical characteristics suited to rearing and harvesting as well as those that can thrive in particular environmental conditions.</p>	<p>Former key areas 3.1 (a) (i) and (ii) combined.</p> <p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>(b) Photosynthesis. Light energy is absorbed by photosynthetic pigments to generate ATP and for photolysis.</p> <p>Absorption spectra of chlorophyll a and b and carotenoids compared to action spectra for photosynthesis. Carotenoids extend the range of wavelengths absorbed and pass the energy to chlorophyll for photosynthesis.</p> <p>Absorbed light energy excites electrons in the pigment molecule. Transfer of these electrons through the electron transport chain releases energy to generate ATP by ATP synthase. Energy is also used for photolysis, in which water is split into oxygen, which is evolved, and hydrogen, which is transferred to the coenzyme NADP.</p> <p>In the carbon fixation stage (Calvin cycle), the enzyme RuBisCO fixes carbon dioxide by attaching it to ribulose bisphosphate (RuBP). The 3-phosphoglycerate (3PG) produced is phosphorylated by ATP and combined with hydrogen from NADPH to form glyceraldehyde-3-phosphate (G3P). G3P is used to regenerate RuBP and for the synthesis of glucose. Glucose may be used as a respiratory substrate, synthesised into starch or cellulose or passed to other biosynthetic pathways.</p>	<p>Light energy not absorbed is transmitted or reflected.</p> <p>Each pigment absorbs a different range of wavelengths of light.</p> <p>These biosynthetic pathways can lead to the formation of a variety of metabolites such as DNA, protein and fat.</p>	<p>Formerly key area 3.1 (b) (i) 'Plant Growth and Productivity'.</p> <p>Minor rewording for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>Reference to electrons being 'high energy' removed.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
		Former key area 3.1 (b) (ii) removed.
<p>2 Plant and animal breeding</p> <p>(a) Plant and animal breeding to improve characteristics to help support sustainable food production.</p>	Breeders develop crops and animals with higher food yields, higher nutritional values, pest and disease resistance and ability to thrive in particular environmental conditions.	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>(b) Plant field trials are carried out in a range of environments to compare the performance of different cultivars or treatments and to evaluate GM crops.</p> <p>In designing field trials account has to be taken of the selection of treatments, the number of replicates and the randomisation of treatments.</p>	The selection of treatments to ensure valid comparisons, the number of replicates to take account of the variability within the sample, and the randomisation of treatments to eliminate bias when measuring treatment effects.	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>
<p>(c) Inbreeding</p> <p>In inbreeding, selected related plants or animals are bred for several generations until the population breeds true to the desired type due to the elimination of heterozygotes.</p>	Analysis of patterns of inheritance in inbreeding using monohybrid crosses.	<p>New key area heading.</p> <p>Reworded for clarification.</p> <p>References to cross- and self-pollinating plants removed.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>A result of inbreeding can be an increase in the frequency of individuals who are homozygous for recessive deleterious alleles. These individuals will do less well at surviving to reproduce. This results in inbreeding depression.</p>		Section on outbreeding removed.
<p>(d) Cross breeding and F₁ hybrids. In animals, individuals from different breeds may produce a new crossbred population with improved characteristics. The two parent breeds can be maintained to produce more crossbred animals showing the improved characteristic.</p> <p>In plants, F₁ hybrids, produced by the crossing of two different inbred lines, create a relatively uniform heterozygous crop. F₁ hybrids often have increased vigour and yield.</p> <p>In inbreeding animals and plants, F₁ hybrids are not usually bred together as the F₂ produced shows too much variation.</p>	<p>New alleles can be introduced to plant and animal lines by crossing a cultivar or breed with an individual with a different, desired genotype.</p> <p>Plants with increased vigour may have increased disease resistance or increased growth rate.</p>	<p>Former key areas 2 (d) and 2 (e) combined.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>Determining F₂ genotype by backcross removed.</p> <p>Testcrosses to identify F₂ genetic variables removed.</p>
<p>(e) Genetic technology. As a result of genome sequencing, organisms with desirable genes can be identified and then used in breeding programmes.</p>	<p>Single genes for desirable characteristics can be inserted into the genomes of crop plants, creating genetically modified plants with improved characteristics.</p>	<p>Formerly key area 3.2 (f).</p> <p>No change to content.</p> <p>Reworded for clarification.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>Breeding programmes can involve crop plants that have been genetically modified using recombinant DNA technology.</p>	<p>Details of recombinant DNA technology techniques in improving crop plants are not required, eg the use of Agrobacterium.</p> <p>Recombinant DNA technology in plant breeding includes insertion of Bt toxin gene into plants for pest resistance, glyphosate resistance gene inserted for herbicide tolerance.</p>	<p>Clarification of required depth of knowledge.</p> <p>Reference to genetic transformation techniques removed.</p>
<p>3 Crop protection</p> <p>(a) Weeds compete with crop plants, while other pests and diseases damage crop plants, all of which reduce productivity.</p> <p>Properties of annual weeds — rapid growth, short lifecycle, high seed output and long-term seed viability.</p> <p>Properties of perennial weeds with competitive adaptations — storage organs and vegetative reproduction.</p> <p>Most of the pests of crop plants are invertebrate animals such as insects, nematode worms and molluscs.</p>		<p>No change to content.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
Plant diseases can be caused by fungi, bacteria or viruses, which are often carried by invertebrates.		
(b) Control of weeds, other pests and diseases by cultural methods.	Ploughing, weeding and crop rotation.	Information on plant protection chemicals moved to 3.3 (c). Clarification of required depth of knowledge.
(c) The advantages of pesticides which are either selective or systemic.	<p>Pesticides include herbicides to kill weeds, fungicides to control fungal diseases, insecticides to kill insect pests, molluscicides to kill mollusc pests and nematicides to kill nematode pests.</p> <p>Selective herbicides have a greater effect on certain plant species (broad leaved weeds).</p> <p>Systemic herbicide spreads through vascular system of plant and prevents regrowth.</p> <p>Systemic insecticides, molluscicides and nematicides spread through the vascular system of plants and kill pests feeding on plants.</p>	<p>Information from former key area 3.3 (b) included here.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>The phrase 'plant protection chemicals' replaced with 'pesticides'.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
Problems with pesticides: toxicity to non-target species, persistence in the environment, bioaccumulation or biomagnification in food chains, producing resistant populations of pests.	<p>Applications of fungicide based on disease forecasts are more effective than treating diseased crops.</p> <p>Bioaccumulation is a build-up of a chemical in an organism. Biomagnification is an increase in the concentration of a chemical moving between trophic levels.</p>	
<p>(d) Control of weeds, other pests and diseases by biological control and integrated pest management.</p> <p>Risks with biological control.</p>	<p>In biological control the control agent is a natural predator, parasite or pathogen of the pest.</p> <p>Integrated pest management is a combination of chemical, biological and cultural control.</p> <p>The control organism may become an invasive species, parasitise, prey on or be a pathogen of other species.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>4 Animal welfare</p> <p>The costs, benefits and ethics of providing different levels of animal welfare in livestock production.</p>	<p>Intensive farming is less ethical than free range farming due to poorer animal welfare.</p> <p>Free range requires more land and is more labour intensive but can be sold at a higher price and animals have a better quality of life.</p>	<p>Formerly key area 3.4 (a).</p> <p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
Behavioural indicators of poor animal welfare are stereotypy, misdirected behaviour, failure in sexual or parental behaviour and altered levels of activity.	Intensive farming often creates conditions of poor animal welfare but is often more cost effective, generating higher profit as costs are low. Very low (apathy) or very high (hysteria) levels of activity.	
		Former key area 3.4 (b) removed.
5 Symbiosis Symbiosis — co-evolved intimate relationships between members of two different species.	Types of symbiotic relationship — parasitism and mutualism. Knowledge of commensalism is not required.	No change to content. Clarification of required depth of knowledge.
(a) Parasitic relationships and transmission. A parasite benefits in terms of energy or nutrients, whereas its host is harmed by the loss of these resources.		No change to content. Minor rewording for clarification.

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>Parasites often have limited metabolism and cannot survive out of contact with a host.</p> <p>Transmission of parasites to new hosts using direct contact, resistant stages and vectors.</p> <p>Some parasitic lifecycles involve intermediate (secondary) hosts to allow them to complete their lifecycle.</p>		
(b) Mutualism. Both mutualistic partner species benefit in an interdependent relationship.		Evolution of mitochondria and chloroplasts removed.
<p>6 Social behaviour</p> <p>(a) Many animals live in social groups and have behaviours that are adapted to group living such as social hierarchy, co-operative hunting and social defence.</p>	<p>Social hierarchy is a rank order within a group of animals consisting of a dominant and subordinate members. In a social hierarchy, dominant individuals carry out ritualistic (threat) displays whilst subordinate animals carry out appeasement behaviour to reduce conflict.</p> <p>Social hierarchies increase the chances of the dominant animal's favourable genes being passed on to offspring. Animals often form alliances in social hierarchies to increase their social status within the group.</p>	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
	<p>Co-operative hunting may benefit subordinate animals as well as dominant ones, as they may gain more food than by foraging alone. Less energy is used per individual. Co-operative hunting enables larger prey to be caught and increases the chance of success.</p> <p>Social defence strategies increase the chance of survival as some individuals can watch for predators whilst others can forage for food. Groups adopt specialised formations when under attack protecting their young.</p>	
<p>(b) Altruism and kin selection and its influence on survival.</p> <p>An altruistic behaviour harms the donor individual but benefits the recipient. Behaviour that appears to be altruistic can be common between a donor and a recipient if they are related (kin).</p> <p>The donor will benefit in kin selection in terms of the increased chances of survival of shared genes in the recipient's offspring or future offspring.</p>	<p>Reciprocal altruism, where the roles of donor and recipient later reverse, often occurs in social animals.</p>	<p>No change to content.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>(c) Social insects and the structure of their society in which only some individuals (queens and drones) contribute reproductively. Most members of the colony are sterile workers who co-operate with close relatives to raise relatives.</p>	<p>Social insects include bees, wasps, ants and termites.</p> <p>Other examples of workers' roles include defending the hive, collecting pollen and carrying out waggle dances to show the direction of food.</p> <p>Sterile workers raise relatives to increase survival of shared genes.</p>	<p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>Ecological importance and evolution of societies of social insects removed.</p> <p>Economic importance removed.</p>
<p>(d) Primate behaviour.</p> <p>Primates have a long period of parental care to allow learning of complex social behaviour.</p> <p>Complex social behaviours support the social hierarchy. This reduces conflict through ritualistic display and appeasement behaviour. Alliances form between individuals, which are often used to increase social status within the group.</p>	<p>Grooming, facial expression, body posture and sexual presentation.</p>	<p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p> <p>Section on the influence of external factors removed.</p>
		<p>Former key area 3.7 (a) removed.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>7 Components of biodiversity Components of biodiversity are genetic diversity, species diversity and ecosystem diversity.</p> <p>Genetic diversity is the number and frequency of all the alleles within a population.</p> <p>Species diversity comprises the number of different species in an ecosystem (the species richness) and the proportion of each species in the ecosystem (the relative abundance).</p> <p>Ecosystem diversity refers to the number of distinct ecosystems within a defined area.</p>	<p>If one population of a species dies out then the species may have lost some of its genetic diversity, and this may limit its ability to adapt to changing conditions.</p> <p>A community with a dominant species has a lower species diversity than one with the same species richness but no particularly dominant species.</p>	<p>New key area heading.</p> <p>Former parts 3.7 (b) (i), (ii) and (iii) combined.</p> <p>Clarification of required depth of knowledge.</p> <p>Effects of degree of isolation and size of habitat islands removed.</p>
<p>8 Threats to biodiversity (a) Exploitation and recovery of populations and the impact on their genetic diversity.</p>	<p>With overexploitation, populations can be reduced to a low level but may still recover. Some species have a naturally low genetic diversity in their population and yet remain viable.</p>	<p>No change to content.</p> <p>Minor rewording for clarification.</p> <p>Clarification of required depth of knowledge.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>The bottleneck effect — small populations may lose the genetic variation necessary to enable evolutionary responses to environmental change.</p>	<p>In small populations, this loss of genetic diversity can be critical for many species, as inbreeding can result in poor reproductive rates.</p>	
<p>(b) Habitat loss, habitat fragments and their impact on species richness.</p> <p>The clearing of habitats has led to habitat fragmentation. Degradation of the edges of habitat fragments results in increased competition between species as the fragment becomes smaller. This may result in a decrease in biodiversity.</p> <p>To remedy widespread habitat fragmentation, isolated fragments can be linked with habitat corridors.</p>	<p>More isolated fragments and smaller fragments exhibit a lower species diversity.</p> <p>The corridors allow movement of animals between fragments increasing access to food and choice of mate. This may lead to recolonisation of small fragments after local extinctions.</p>	<p>No change to content.</p> <p>Reworded for clarification.</p> <p>Clarification of required depth of knowledge.</p>
<p>(c) Introduced, naturalised and invasive species and their impact on native populations.</p> <p>Introduced (non-native) species are those that humans have moved either intentionally or accidentally to new geographic locations.</p>		<p>No change to content.</p>

Key areas	Depth of knowledge required	Changes to content
Sustainability and interdependence		
<p>Those that become established within wild communities are termed naturalised species. Invasive species are naturalised species that spread rapidly and eliminate native species therefore reducing species diversity. Invasive species may well be free of the predators, parasites, pathogens and competitors that limit their population in their native habitat. Invasive species may prey on native species, out-compete them for resources or hybridise with them.</p>		<p>Reworded for clarification.</p>
		<p>Former key area 3.8 (d) removed.</p>