			Section A
			Answer all questions.
1	Anir	nals tha	at possess hair are classified within the phylum Chordata and class Mammalia.
	(a)	Define 1. Bid us 2. It u int ma 3. Ph	 biological classification and explain how classification relates to phylogeny.[3] clogical classification groups organisms based on <u>overall morphological similarities</u> and ually does not <u>consider their evolutionary history;</u> uses a <u>naming system</u> where each organism is given a <u>binomial name</u>* and grouped o a <u>domain, kingdom, phylum, class, order, family, genus and species</u> in a <u>hierarchical anner;</u> hylogeny <u>also groups organisms</u> but they are grouped based on their <u>evolutionary history</u>
		/ <u>a</u> 4. <u>It</u> <u>ph</u> tre	ncestor-descendent relationships using molecular data and; does not rank organisms but instead assigns each organism a position on a nylogenetic tree * based on its evolutionary relationship with other organisms on the e;
	(b)	(i)	Epithelial stem cells and melanocyte stem cells are examples of a particular type of stem cell. State two key features of this type of stem cell. [2]
			 They are <u>multipotent</u>* stem cells that can differentiate into a <u>limited range of</u> specialized cell types in the presence of the <u>correct molecular signals</u>;
			 They are able to undergo <u>extensive proliferation</u> and <u>self-renewal</u> by <u>mitosis</u>* in the presence of the <u>correct molecular signals</u> and hence <u>give rise to a constant pool of</u> <u>cells</u>;
		(ii)	Discuss how the presence of stem cells within the skin and the presence of hairs covering the skin influence the harmful effects of high levels of ultraviolet light on the health of mammals. [4]
			 The presence of hairs covering the skin play a protective role as they reduce exposure of the skin by blocking the harmful ultraviolet radiation; Since stem cells are near the skin surface, the high levels of ultra violet light can damage the DNA is stem cells as; the ionizing radiation can cause the DNA is the stem cells in the skin to mutate and if these mutations are in genes that regulate cell cycle checkpoints/proto oncogenes/ tumour suppressor genes; it could lead to uncontrolled cell division/ tumour formation and increase the chances of an individual getting cancer; especially because stem cells already have telomerase activity;



	(ii)	In cats with agouti hairs, ASIP is secreted at intervals during hair growth.
		With reference to Table 1.1 and the information provided, explain how this interrupted pattern of ASIP secretion leads to striped agouti hairs in cats.[4]
		 During intervals when <u>ASIP is not secreted</u>, <u>α-MSH</u> binds to MC1R (a GPCR), causing the MC1R to undergo a <u>conformation change in its intracellular domain</u> allowing <u>G-protein to bind to it</u>; and
		2. <u>G-protein is activated</u> when it <u>displaces its attached GDP</u> for GTP;
		 Activated G-protein will <u>translocate along membrane</u> and <u>bind to enzyme</u> adenylyl cyclase and phosphorylate it, thus activating it;
		 Adenylyl cyclase will catalyze conversion of <u>ATP to cAMP</u>, which will trigger a phosphorylation cascade that will cause <u>black eumelanin pigment to be</u> <u>produced</u>, making the hair black;
		 However, during intervals when <u>ASIP is secreted</u>, ASIP will prevent <u>α-MSH</u> from binding to MC1R/ ASIP will preferentially bind to MC1R;
		 Thus MC1R will not undergo a conformational change that allows G-protein to bind to it and so the <u>G protein will not be activated</u>, adenylyl cyclase will not activated, cAMP is not produced and so yellow phaeomelanin pigment is produced making the hair yellow;
		Hence the interrupted pattern of ASIP secretion, will lead to the striped yellow and black agouti hairs in cats.

Table 1.2 number of individuals of each genotype number of fur colour species individuals genotype at ASIP locus genotype at MC1R locus phenotype in sample $\Delta 2/\Delta 2$ $\Delta 2/+$ +/+ A15/A15 $\Delta 15/+$ +/+ 26 0 F. catus agouti 15 11 0 0 26 F. catus black 57 0 0 0 0 57 57 P. onca agouti 36 0 0 36 0 0 36 P. onca black 10 0 0 1 9 10 0 key $\Delta 2$ = allele with 2 base pair deletion of nucleotides 123–124 of the ASIP gene + = normal allele of the ASIP or MC1R gene $\Delta 15$ = allele with 15 base pair deletion of nucleotides 301–315 of the MC1R gene (d) (i) Use Table 1.2 to identify: a recessive allele: $\wedge 2$ a dominant allele : 15 an allele containing a frameshift mutation $\triangle 2$ [3] (ii) With reference to Table 1.1, Table 1.2 and the information provided, explain why melanic (black) individuals of *F. catus* have hairs that do not contain any phaeomelanin. [4] 1. Melanic *F.catus* has 2 copies of $\triangle 2$ and hence has 2 copies of the mutated ASIP alleles; 2. A 2 base pair deletion in the gene will result in a frame shift in the DNA resulting in a change the sequence of *codons** in the mRNA sequence; 3. which will cause a change in the sequence of amino acids in the polypeptide which in turn will affect the R group interactions between the amino acids in the tertiary structure of the ASIP protein, resulting in a change in its 3D conformation; A: truncated non-functional polypeptide 4. As a result, the non-functional ASIP (ligand) cannot bind to the MC1R (receptor) on the melanocyte cell as it is no longer *complementary** in shape to it; 5. Thus and <u>cAMP (second messenger) production will continue and black eumelanin</u> is produced in the melanocyte cell; Thus the melanic individuals have hairs that do not contain any phaeomelanin; (iii) Scientists have concluded that the black fur colour phenotype found in several cat species has evolved more than once. Use information in Table 1.2 to justify this conclusion.[1] Since cats with the same phenotype, that is, having black fur, have 2 different mutations in 2 different gene loci, either in the ASIP locus or the MC1R locus, it shows that black fur colour phenotype in cat species evolved more than once;

4

(e)	(i)	Explain Fig. 1.1.	now temperature va [2]	ariation of the skin co	ontri	butes to the phe	enoty	pe of the cat in
		 Since tempe hence The <u>c</u> tyrosi the pr 	the <u>ears, face, pay</u> erature sensitive tyr e <u>black pigment car</u> ther parts of the bo nase enzyme starts roduction of pigmen	vs and tail are the <u>co</u> osinase enzyme <u>do</u> to be produced in the dy are warmer abov to denature, lowering the the hairs there;	<u>oole</u> <u>es n</u> haii <mark>/e 3(</mark> ng e	r parts of the bo ot denature in the <u>'s</u> in those areas <mark>)°C</mark> and the <u>tem</u> enzyme activity a	<u>dy</u> of nose s; pera and h	the cat, the <u>areas</u> and <u>ture sensitive</u> ence <u>reducing</u>
	(ii)	Draw a g cross. Insert th for the a genetic o	genetic diagram to s e symbols A and a lleles of the tyrosina diagram. [4]	show the predicted of for the alleles of the ase gene in the key	AS to s	colours of the o <i>IP</i> gene and the ymbols table, be	ffspri sym efore	ng of this bols H and h completing the
		$\Delta 2$ allel	e of the ASIP gene			а		
		normal	allele of the ASIP of	jene		Α		
		Himala	yan allele of the tyre	osinase gene		h		
		non-Hir	malayan allele of the	e tyrosinase gene		н		
gen Par Par	etic dia ental pl ental g	agram sho henotype enotype	wing predicted coat Himalayan cat w black extremities aahh	t colours: /ith X S	Ago	outi, Non-Himala AaHl	iyan (cat
Gar	netes		ah		(A	H Ah aH		h
Ga	ametes		AH	Ah		aH		ah
	ah)	AaHh	Aahh		aaHh		aahh
			Agouti, Non- Himalayan cat	Himalayan cat with Agouti extremities	BI Hi	ack, Non- malayan cat	Him with extr	alayan cat black emities
Offs	spring g	genotype:	AaHh	Aahh		aaHh		aahh
Offs	spring p	henotype	Agouti, Non- Himalayan cat	Himalayan cat with Agouti	B Hi	lack Non- malayan cat	H w	imalayan cat ith black
Offs ratio	spring p	henotypic	: 1 :	extremities 1 :		1	: E	tremities 1

1m 1 1m 1 1m 1 1m 1	for corr for cor for corr for corr	ect parental phenoty rect gametes which a ect offspring genotyp ect phenotypic ratio ;	be and are circl e and p	genotype ; led ; phenotype ;				
	(iii)	Put a tick (\checkmark) in one Give a reason for yo	box to	o indicate whethe	er or not this cross i	is a test (cros	S.
 		test cross	✓		not a test cross			
		reason : The hetero both the mutant AS	zygous IP and	s cat was <u>crosse</u> heat sensitive ty	<u>ed with a homozygo</u> vrosinase gene loci	<u>us reces</u>	sive	<u>cat</u> with
	(iv)	Name a statistical te genetic diagram by your expected resul	est that compa ts.[1]	would allow you ring the observe	u to test the assumped results of many o	otions yo crosses c	ou m of thi	ade in your s type with
		Chi-square test						
 								[Total: 30]

2	(a)	Descr meas	ibe and explain the relationship shown in Fig. 2.1 between the number of cases of les and the percentage uptake of vaccine.[4]
		 Fr th Af of Ar <l< td=""><td>om 1969 to 1996, as the percentage of uptake of vaccine increases from 32% to 94%, e number of measles cases decreases from 320 000 to 0 cases; ter 1996, as the percentage of uptake of vaccine remained high above 80%, the number measles cases were 0; ntigen presenting cells (APCs) macrophages / dendritic cells (A: antigen processing cells) ke up the measles vaccine <u>by phagocytosis</u>, process <u>antigen</u> and <u>present</u> it as a <u>aptide:MHC complex</u>; PC activates naïve T cell which will undergo <u>clonal expansion and differentiation</u> to form eleper T cells, cytotoxic T cells and memory T cells; <u>helper cells</u>* bind to secrete cytokines that <u>activate specific naïve B cells*</u> to undergo onal expansion and differentiation and form antibody-secreting plasma cells and memory cells; <u>emory B and T cells</u>* when <u>exposed to virus</u>, will recognize it and mount a <u>faster</u> and ronger and longer lasting <u>secondary immune response</u>; <u>becific antigen binding site</u> of antibody <u>binds to specific epitope on antigen</u> of coming virus; revent/block viral glycoproteins from binding to host cell receptors, resulting in <u>autralization</u>*; revents attachment and subsequent infection of cells, t<u>hus reducing the number of</u> <u>ises of infection</u>;</td></l<>	om 1969 to 1996, as the percentage of uptake of vaccine increases from 32% to 94%, e number of measles cases decreases from 320 000 to 0 cases; ter 1996, as the percentage of uptake of vaccine remained high above 80%, the number measles cases were 0; ntigen presenting cells (APCs) macrophages / dendritic cells (A: antigen processing cells) ke up the measles vaccine <u>by phagocytosis</u> , process <u>antigen</u> and <u>present</u> it as a <u>aptide:MHC complex</u> ; PC activates naïve T cell which will undergo <u>clonal expansion and differentiation</u> to form eleper T cells, cytotoxic T cells and memory T cells; <u>helper cells</u> * bind to secrete cytokines that <u>activate specific naïve B cells*</u> to undergo onal expansion and differentiation and form antibody-secreting plasma cells and memory cells; <u>emory B and T cells</u> * when <u>exposed to virus</u> , will recognize it and mount a <u>faster</u> and ronger and longer lasting <u>secondary immune response</u> ; <u>becific antigen binding site</u> of antibody <u>binds to specific epitope on antigen</u> of coming virus; revent/block viral glycoproteins from binding to host cell receptors, resulting in <u>autralization</u> *; revents attachment and subsequent infection of cells, t <u>hus reducing the number of</u> <u>ises of infection</u> ;
	(b)	lf the thresh vaccir	percentage of people who are immune to a disease exceeds the herd immunity hold, the disease can no longer persist in the population. Assuming 100% efficacy of the herd immunity threshold is calculated as: $100 \times (1 - 1/R)$
		(i)	Calculate the herd immunity threshold for measles, if $R_0 = 11$.
			herd immunity threshold =% [1] 90.9 or 91%
		(ii)	Use Fig. 2.1 and your answer to (b)(i) to identify the years when the herd immunity threshold was achieved or exceeded in this part of the United Kingdom.[1]
			from 1992 to 1999 inclusive

(c)	The m substi to be 2	nutation rate in the measles virus haemagglutinin gene is estimated to be 6.0 x 10^{-4} tutions per nucleotide per year. The mutation rate in human nuclear DNA is estimated 2.5 x 10^{-8} substitutions per nucleotide per generation time of 20 years.
	(i)	Suggest reasons for the difference in mutation rate of the measles virus haemagglutinin gene and the mutation rate of human nuclear DNA.[2]
		 Measle virus has a single-stranded RNA genome unlike human DNA that is double stranded; No backup copy to do correction and hence has a higher mutation rate; OR (RNA-dependent/ viral) <u>RNA polymerase lacks proof-reading ability unlike human RNA polymerase that has proof reading ability;</u> more errors occur in viral genome <u>replication;</u> RNA genome more reactive (due to the 2'OH group) than DNA genome; R: unstable <u>Higher frequency of mutations of the RNA</u>* genome leading to the <u>changes in the RNA sequence</u> Note: It is important to make comparative statements. Use of words like more and higher are important
	<u>(ii)</u>	 Explain why knowledge of mutation rates is useful in reconstructing phylogenies.[2] <u>Rate of mutation/time taken for mutation to accumulate</u> in virus types is <u>constant</u>; from the <u>number of nucleotide differences between different virus strains</u>, it is possible to <u>distinguish between different virus strains</u> /infer evolutionary relationships/degree of genetic divergence between different virus strains or <u>infer common ancestor</u> of virus strains; from the <u>number of nucleotide differences</u>, it is possible to <u>infer the time when a certain strain virus emerged or share common ancestor</u>; Reject time of speciation because viruses are not able to form new species. A: if not specifically reference to virus
		[Total: 10]

3	(a)	Name growt	e these two products and explain how the presence of each of these could help coral h.[4]
		1. 2. 3. 4. 5.	 Oxygen* is produced by the algae during the light dependent reaction of photosynthesis; It is used by the corals for <u>aerobic respiration</u> as the <u>final electron acceptor</u>* of the <u>electron transport chain*</u> during <u>oxidative phosphorylation</u>*; <u>Glucose</u>* is produced by algae during the <u>light independent reaction</u> of photosynthesis; It is the main <u>substrate</u> of <u>respiration</u> by the corals; To produce energy in the form of <u>ATP</u>* for metabolism and growth;
	(b)	(i)	Calculate the overall mean percentage changes in coral cover per year in area 1 and area 2 from 1985 to 2012. State which area shows the greater overall mean percentage change in coral cover per year.
			overall mean percentage change in area $1 = (+2.07 - 0.77 - 1.05 - 0.36)\% = -0.11\%$ overall mean percentage change in area $2 = (+2.34 - 1.59 - 1.75 - 0.04)\% = -1.04\%$
			overall mean percentage change in coral cover per year in area 1 =0.11%
			overall mean percentage change in coral cover per year in area 2 =1.04%
			area with greater overall mean percentage change in coral cover per year =area 2[2]
		(ii)	Use the data in Table 3.1 to evaluate whether or not the changes in coral cover in these two areas of the Great Barrier Reef from 1985 to 2012 can be attributed to global climate change.[4]
			 Decrease can be attributed to climate change because Coral bleaching is a result of <u>increase in sea water temperatures</u> which cause <u>zooxanthellae to be expelled</u> from corals, eventually leading to death of coral; Mean percentage decrease due to coral bleaching was <u>0.36% and 0.04%</u> in <u>area 1 and area 2</u> respectively; Tropical storms are <u>more severe due to increased temperatures</u> from climate change; Mean percentage decrease due to tropical storms <u>1.05% and 1.75%</u> in <u>area 1 and area 2</u> respectively; Decrease should not be attributed to climate change because
			 There were <u>only 2 sites</u> studied, both in the Great Barrier Reef; <u>Changes may be due to localised environmental conditions</u> and not global climate change; Data is represented as a <u>mean value over 27 years</u>, so <u>unable to see changes</u> <u>on a yearly basis</u> to make an informed conclusion;

		Section B Answer one question in this section	
-			.
4	(a)	Outline the structural features and genomic organisation of bacteria.	[15]
		1. Bacteria cells have simple internal structures with no membrane-bound	
		organelles*.	
		2. The <u>bacterial chromosome</u> is a <u>double-stranded, circular DNA molecule</u> *;	
		3. that makes up a dense region within the cell called the <i>nucleoid</i> [*] region;	
		4. Some bacteria cells may contain <u>extrachromosomal</u> <u>double-stranded small</u>	
		5. It contains 70S ribosomes * needed for protein synthesis:	
		6. Nutrients and chemical reserves may be stored in the cytoplasm in the form of	
		granules, e.g. glycogen granules, lipid granules, etc;	
		7. The bacterial cytosol is enclosed by <u>cell surface membrane</u> * made up of a	
		phospholipid bilayer * with proteins embedded in carrying out specific functions	
		such as transport, ATP synthesis, etc;	
		6. The peptidogrycan cen wan conters rightly to the bacterial cen and protects it from osmotic lysis:	
		9. Gram-positive bacteria have a cell wall with a thick peptidoglycan layer;	
		10. Gram-negative bacteria have a cell wall with a thin peptidoglycan layer;	
		11. Some bacteria may have a layer of polysaccharides known as glycocalyx to the	
		exterior of the cell wall;	
		12. If the <u>diveccality layer</u> is a <u>distinct layer</u> , it is known as a <u>capsule</u> ;	
		14. Some bacteria may have short, bristle-like fibres extending from the cell surface	
		called <i>fimbriae</i> *;	
		15. Fimbriae may be evenly distributed over the entire cell surface or at the poles of the cells;	
		16. Other bacterial appendages include <i>pili</i> *, which are longer, hollow hair-like structures found on the cell surface:	
		17. They occur in fewer numbers compared to fimbriae and may have specialized	
		functions such as the sex pilus during conjugation;	
		 Some bacteria cells may have a <u>long, hollow cylindrical protein thread</u> called the <u>flagella</u>*; 	
		19. Flagella are usually <u>used by bacteria cells for motility</u> as it helps <u>propel the</u> <u>bacteria by rotation</u> ;	
		(B) Genomic organization of bacteria	
		20. The bacterial chromosome is small in size about $10^4 - 10^7$ base pairs;	
		21. It comprises of a single double-stranded, circular DNA molecule [*] ;	
		<u>histone-like proteins,</u> followed by further <u>supercoiling</u> to form highly condensed DNA	
		23. that makes up a dense region within the cell called the <i>nucleoid</i> * region;	
		24. The genome is simple with a <i>single origin of replication</i> *;	
		25. where genes with related functions or belonging to the same metabolic pathway	
		are <u>organized together</u> in an <u>operon*</u> under the <u>control of a single promoter</u> *;	
		26. Bacterium being prokaryotic, has <u>no intron sequences</u> in its chromosome;	
		circular DNA molecules called plasmids *.	
		28. There may be more than 1 copy of the same plasmid in the bacterial cell:	
		29. The plasmid may contain genes which may confer advantages for the bacteria living in stressful environments, e.g. antibiotic resistance genes;	
		QWC: Address both (A) and (B)	

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Point of comparison	prokaryote	eukaryote
1. Method of obtaining genetic variation	1A. Mainly by horizontal gene transfer;	1B. Mainly by meiosis and sexual reproduction;
2. Methods of genetic variation	2A1. <u>Conjugation</u> whereby the <u>F plasmids</u> to transferred from host to the recipient via a mating bridge;	2B1. Meiosis: During <u>crossing over between</u> <u>non-sister chromatids of</u> <u>homologous</u> <u>chromosomes</u> * in <u>prophase</u> <u>1</u> results in new
	2A2. <u>Specialised</u> <u>transduction</u> * whereby a <u>temperate phage takes up an</u> <u>adjacent bacterial DNA of the</u> <u>provirus</u> when it undergoes the lytic phase and phage infects another bacteria. This is transferred to another bacteria via homologous recombination;	combinations of alleles on chromatids. (& eventually a variety of offspring) 2B2. <u>Independent</u> <u>assortment of homologous</u> <u>chromosomes at the</u> <u>metaphase plate</u> & their subsequent separation during metaphase I & anaphase I respectively &
	2A3. <u>Transformation*</u> whereby fragments of naked DNA are <u>taken in</u> by a competent bacteria cell and introduced into the <u>bacterial chromosome by</u> crossing over of homologous regions;	2B2. <u>Random orientation of</u> <u>non-identical sister</u> <u>chromatids of each</u> <u>chromosome at the</u> <u>metaphase plate</u> & their subsequent separation during metaphase II and anaphase II respectively
		2B3. This results in <u>gametes</u> with <u>different combinations</u> <u>of maternal & paternal</u> <u>chromosomes</u> . (& eventually in a variety of offspring)
		3. <u>Random fusion of</u> <u>gamete occurs during sexual</u> <u>reproduction</u> /fertilisation results in offspring with a variety of genotypes & possibly phenotypes (& hence a variety of offspring);
4. Mutations	As there is only 1 chromosome, there will not be mutations that involve more than 1	As there are more than 1 chromosome, non- disjunction, translocation

	disjunction, polyploidy and translocation Similarities: 5. Mutations are a source of genetic variation for both prokaryotes and eukaryotes; 6. Mutations such as point mutations and gene mutations such as inversion, substitution, deletion and addition can occur in both; 7. Role of mutations in both can be significant as novel phenotypes can be selected for during natural selection hence increasing the frequency of alleles coding for the	
	favourable phenotype in the subsequent generation;	
	8. QWC: Mention both similarities and differences.	
	[Tot	al: 25]
5 (a)	 Outline the processes that occur in aerobic respiration in eukaryotic cells. 1. Aerobic respiration occurs when organic food substances such as carbohydrates, fats and proteins are oxidised to <u>provide energy</u>, protons and electrons that will ultimately be used to generate ATP; 2. Oxvgen is required for the complete oxidation of these organic food substances and <u>ATP is produced</u> in the process; (A) Glycolysis 3. Glycolysis* occurs in the <u>cytoplasm</u> of the eukaryotic cell; 4. Phosphorylation of glucose involves the initial investment of 2 ATP molecules which phosphorylate the two ends of glucose molecule forming fructose-1,6-bisphosphate*; 5. Phosphoructokinase* is the enzyme involved in the second phosphorylation step; 6. The phosphorylated <u>6C sugar splits into two 3C sugar phosphates;</u> 7. Each fructose molecule ultimately gives rise to <u>two molecules of glyceraldehyde-3-phosphate (G3P)</u>; 8. Glyceraldehyde-3-phosphate undergoes <u>oxidation by dehydrogenation</u>* to form 1,3-bisphosphoglycerate; 9. NAD* is produced by <u>substrate level phosphorylation</u>* of 1,3-bisphosphoglycerate forming <u>pyruvate</u>*; 11. Since each glyceraldehyde-3-phosphate yields 2 ATP and 1 NADH, <u>each glucose molecule yields 2 ATP, 2 pyruvate and 2 NADH</u> taking into account the 2 ATP as an initial investment; (B) (B) Link reaction 12. Link reaction* occurs in the <u>matrix* of the mitochondria;</u> 13. Pyruvate formed from glycolysis enters the mitochondria; 14. Pyruvate undergoes <u>oxidative decarboxylation</u>* where it combines with coenzyme A to form <u>acetyl CoA(2C)*;</u> 15. Carbon dioxide* is released and <u>NAD*</u>; is reduced to <u>NADH</u>* in the process; 16. Each glucose molecule produces 2 pyruvate which undergoes the link reaction to <u>produce 2 acetyl CoA, 2 CO₂ and 2 NADH;</u> 	[15]

(C)	(C) Krebs cycle
	17. Krebs cycle* occurs in the matrix* of the mitochondria and when oxygen is
	present;
	 <u>Acetyl CoA (2C)</u> combines with oxaloacetate (4C)* to form citrate (6C)*;
	19. Citrate is <u>decarboxylated</u> and <u>dehydrogenated</u> to form <u><i>α-ketoglutarate (5C)</i>*</u>
	and <u>NADH</u> .
	20. Each decarboxylation step results in a loss of carbon in the form of <u>a carbon</u>
	<u>dioxide;</u>
	debudrogenetion stops to viold 2 NADH 1 EADH and 1 CO :
	22 High energy electrons originally from the glucose molecule have now been
-	transferred to electron carriers NAD ⁺ and FAD ⁻
	23. NAD ⁺ + 2H ⁺ + 2e ⁻ \rightarrow NADH + H ⁺ (or reduced NAD)
	$FAD + 2H^+ + 2e^- \rightarrow FADH_2$ (or reduced FAD)
	24. 1 ATP* is also produced through substrate-level phosphorylation* during this
	regeneration process;
	25. Altogether <u>1 molecule of glucose will yield 6 NADH, 2 FADH₂ & 2 ATP</u> through
	the Krebs cycle. The coenzymes with their reducing power will next be
	transported to the electron transport chain where the bulk of ATP is generated.
(D)	Oxidative phosphorylation
	Oxidative phosphorylation <u>occurs on the</u> folds of the inner membrane of mitophondria
	MILOCHONDHA = <u>Cristae</u> ;
	 <u>INADE dollates electrons to</u> first electron carrier or <u>the electron transport</u> <u>chain*</u> found on the cristae:
	 The electron carriers alternate between reduced and oxidized states as they
	accept and donate electrons:
	Each successive carrier has a higher electronegativity:
	• Each FADH ₂ donates electrons to the chain at a lower level and hence
	generates 2 ATP compared to 3 ATP generated by NADH;
	• Electron transport through the series of carriers is coupled to the active
	pumping of H ⁺ into the intermembrane space. This generates a proton
	gradient*/ proton motive force* across the mitochondria membrane;
2	26. The proton motive force is used to phosphorylate ADP. As protons diffuse back
	into the matrix via the ATP synthase complex, ADP is phosphorylated to ATP *.
	27. This process is known as <u>chemiosmosis</u> *;
	• and produces 90% of the ATP during aerobic respiration.
	 With <u>oxygen</u>[*] as the <i>final electron acceptor</i>[*] having the highest
	• $2e^{-} + 2\pi^{2} + 1/2O_{2} \neq \pi_{2}O_{2}$
ວາທ	C: Must have at least two points from (A) (B) (C) and (D)
QVV	

Fea	ature of comparison	ATP generation in chloroplasts	ATP generation in mitochondria
1.	Process(es) in which ATP is generated	Photophosphorylation* in the light-dependent reaction	Substrate-level phosphorylation* (Krebs cycle) and oxidative phosphorylation* in aerobic respiration
2.	Site of ATP formation	Takes place in/at the <u>stroma*/thylakoid</u> membrane* of chloroplast	Takes place in/at the <u>matrix*/intermembrane</u> space* of mitochondria
3.	Source of energy	Energy for synthesis of ATP comes ultimately from <u>light;</u>	Energy for the synthesis of ATP comes from the <u>oxidation of glucose</u> which stores chemical energy;
4.	Source of electrons which travels down the electron transport chain	Water is the electron donor in the non-cyclic pathway while Photosystem I* is the electron donor in the cyclic pathway;	[oxidative phosphorylation only]: <u>NADH and FADH</u> 2 are the electron donors to the first electron carrier of ETC;
5.	Location of proton reservoir which supplies protons that diffuse down ATP synthase	Protons are pumped from stroma across the thylakoid membrane, into the <u>thylakoid space</u> * to establish a proton gradient for ATP synthesis;	[oxidative phosphorylation only]: Protons are pumped from mitochondrial matrix across the inner membrane, into the <u>intermembrane space*</u> to establish a proton gradient for ATP synthesis;
6.	Direction of proton flow through ATP synthase	Protons diffuse from <u>thylakoid space*</u> to <u>stroma*</u>	Protons diffuse from <u>intermembrane space</u> * to <u>matrix</u> *
7.	Type of phosphate	Inorganic phosphate	[substrate-level phosphorylation only]: phosphate derived from respiratory intermediates/substrate
8.	By-product	Splitting of water produces oxygen as by- product during non-cyclic pathway;	[oxidative phosphorylation only]: <u>Water is produced</u> as by-product when oxygen combines electron and proton at the end of ETC;
9.	Energy conversion	Light energy is converted to chemical energy in the process;	Chemical energy (from glucose) is converted to chemical energy (in the form of ATP) in the process;

 Similarities: 1. Both involve the transfer of electrons down electron carriers* of increasing electronegativity in an electron transport chain*; 2. In both cases, energy release from the transfer of electrons down the electron transport chain* is coupled to the pumping* of H*; 3. Both involve the generation of a proton gradient* across a membrane; 4. Both involve <u>facilitated diffusion* of H* through the hydrophilic pore* of ATP synthase* down its proton gradient;</u> 5. In both, <u>ATP* is produced from ADP* and inorganic phosphate/Pi</u> * via chemiosmosis*; QWC: essays need to include at least one similarity and one difference. 	
[Total: 2	5]