# 2007 'A' Level H2 Biology Mark Scheme

# PAPER 1 (MCQ)

1	С
2	B
3	B
4	С
5	В

6	D
7	В
8	С
9	В
10	С

11	D
12	Α
13	С
14	Α
15	Α

16	D
17	С
18	С
19	D
20	C

21	В
22	С
23	D
24	Α
25	D

26	Α
27	С
28	D
29	D
30	С

31	В
32	D
33	Α
34	С
35	D

36	Α
37	С
38	Α
39	D
40	D

## PAPER 2 (CORE)

### **QUESTION 1**

#### (a)

A – viral RNA

- B reverse transcriptase
- C capsid protein

### (b)

- 1 host cell surface membrane surrounds virus particle / (viral) RNA and protein;
- 2 ref. synthesis of surface glycoprotein;
- 3 budding; *R exocytosis*

(C)

(i)

- 1 block protease active site so that substrate unable to bind;
- 2 viral polyprotein not cut / structural / enzyme protein not formed;
- 3 (newly formed) virus unable to infect further cells;

(ii)

- similar shape to polyprotein / complementary shape to active site;
- 2 inhibitor not hydrolysed so remains in active site;

# **QUESTION 2**

(a)

- 1 fluid refers to the phospholipids and proteins being free to move;
- 2 phospholipids also being able to move within a layer, transversely;
- 3 mosaic refers to the proteins embedded in the phospholipid bilayer, where they are scattered;

(b)

(i)

- 1 glucose is a polar molecule and ions are charged;
- 2 unable to pass through hydrophobic region within the membrane;
- 3 ref. to specificity of transport proteins;

#### (ii)

- high levels of ATP close ATP-sensitive potassium channels;
- 2 build up of potassium ions inside cell;
- 3 depolarize membrane;

## (iii)

- 1 insulin release involved exocytosis and therefore did not require a protein channel;
- 2 insulin is a protein;
- 3 too large to pass through membrane;
- 4 ref. to channel large enough to allow insulin through would allow other molecules through;

## (C)

- 1 vesicles containing insulin move to cell surface membrane;
- 2 membrane of vesicle fuses with cell surface membrane;
- 3 vesicle contents emptied by exocytosis;

## **QUESTION 3**

(a)

- 1 DNA strands separate by hydrogen bonds breaking;
- 2 both strands act as templates;
- 3 each new molecule of DNA contains one new and one original strand of DNA;

## (b)

- 1 ref. to shape of active site of DNA polymerase;
- 2 needs 3'OH to initiate action of DNA polymease;
- 3 primers add only at 3';

(c)

- 1 DNA in bacteria occurs in a loop while human DNA is linear;
- 2 bacteria DNA is naked while human DNA is arranged in nucleosomes wrapped around histones;
- 3 ref. to introns only in human DNA;
- 4 ref. to telomeres only in human chromosomes;
- 5 ref. to centromeres only in human chromosomes;
- 6 ref. to operons only in bacteria;

(i)

- 1 much longer DNA molecule;
- 2 DNA packaging in nucleosomes with DNA wrapped around histones
- 3 more non-coding DNA in human chromosomes;

## (ii)

1 multiple sites of replication (on each chromosome);

<sup>(</sup>d)

### (a)

- (inherited) change in DNA/nucleotide sequence/base sequence;
- 2 base substitution;
- 3 change in codon/triplet for amino acid/change in R group;
- 4 base addition/deletion;
- 5 ref. to frameshift or described;

## (b)

- 1 (if addition/deletion resulting in) frameshift;
- 2 protein coded for no longer synthesized/non-functional;
- 3 (if substitution resulting in) amino acid changed;
- 4 change 3D/tertiary structure of protein/different R group;
- 5 ref. change in binding site ref. to active site is neutral

## (C)

- both copies of a tumor suppressor gene must be mutated (before a person will develop cancer);
- 2 ref. to effect of mutation being masked/ref. to normal allele being dominant/remaining functional copy of gene;
- 3 sufficient quantities of proteins synthesized to be effective;

# (d)

- specific shape of binding site of p53 protein;
- 2 corresponds to/complimentary to shape, part of DNA molecule;
- 3 different sequences of bases have different shapes;
- 4 ref. to major and minor grooves on DNA;

## (e)

- 1 change in amino acid in region away from binding area;
- 2 may not affect shape or binding properties;
- 3 ref. idea some amino acids coded for by more than 1 codon/triplet of bases/ref. genetic code/some codons degenerate;
- 4 change in base/codon may not change amino acid/neutral;

<b>QUESTION</b> (a) RRPP RRPp RrPP RrPP	15			
<b>(b)</b> parents gametes	RRpp x rrP Rp x rp	P;		
F1 gametes	RrPp; RrPp x RrPp RP, Rp, rP, rp		RP, Rp, rP, rp;	
F2	walnut RRPP RRPp x 2 RrPp x 4 Rrpp x 2	rose RRp Rrpp Rrpp	pea rrPP rrPp rrPp	single ;; rrpp

(allow on Punnett square, all genotypes must be shown, correspond to correct phenotype)

(c) RrPP x RRpp RRPP x Rrpp RRPP x RRpp

5

#### (a)

P nucleus / euchromatin / nucleoplasm / chromatin;

**R:** heterochromatin

- Q nuclear envelope;
- R cell surface membrane;

## (b)

- site of cellular respiration / oxidative phosphorylation / aerobic respiration;
- 2 synthesis / production / release of ATP;
- 3 ref. specific use of ATP (e.g. active transport / vesicle transport / mitosis / macromolecule synthesis, incuding DNA replication, transcription and translation);
- 4 ref. ATP produced by mitochondria released into cytosol while ATP in chloroplasts not;

# (c)

- (i)
- 1 double membrane;
- 2 highly folded / large surface area of inner membrane;
- 3 presence of ETC / proton pumps;
- 4 ATP synthase;
- 5 70S ribosomes;
- 6 DNA loops;

## (ii)

- 1 three membranes / double outer membrane in chloroplasts;
- 2 ref. position of stalked particles / chloroplasts "inside out"
- 3 starch grains;
- 4 stack of thylakoid membranes / grana;
- 5 ref. to thylakoids once;
- 6 no cristae in chloroplasts;
- 7 contain pigments / chlorophyll; **A**: other photosynthetic pigments

## (d)

- oxidative phosphorylation: final e<sup>-</sup> (and H<sup>+</sup>) acceptor / combines with e<sup>-</sup> and H<sup>+</sup> to form water;
- 2 photophosphorylation: (waste) produce / from photolysis

(a) (dot correct answers then ticks bottom right of page)

animal ; phylum ; class ; order ; Dicrurus ; species ; 6 correct = 4

- 5 correct = 3
- 4 correct = 2
- 3 correct = 1

(b)

- 1 ref. inherited variation/variation in alleles;
- 2 ref. natural selection;
- 3 differences in selection pressures in different habitats;
- 4 ref. to survive to pass on alleles;
- 5 geographical isolation / ref. allopatric;
- 6 idea that broadleaved woodland not continuous;
- 7 ref. islands;
- 8 prevents interbreeding / gene flow;
- 9 ref. to genetic drift/founder effect;

- 10 similar (morphological/anatomical/physiological) features;
- 11 capable of interbreeding;
- 12 to produce fertile offspring;

#### (a)

### <u>Cellulose</u>

- 1 a polysaccharide;
- 2 (chain of)  $\beta$ /beta glucose;
- 3 <u>1-4</u> links; (glycosidic bonds not as important as linking bonds)
- 4 ref. to microfibrils;
- 5 ref. to bundles of microfibrils/macrofibrils/fibres;

### <u>Collagen</u>

- 1 a <u>fibrous</u> protein;
- 2 (chain of) amino acids;
- 3 ref. to peptide bonds;
- 4 ref. to each chain helical;
- 5 triple helix/tropocollagen;
- 6 ref. to role/presence of glycine;
- 7 ref. to staggered ends;
- 8 ref. to H bonds; A once in either

# (b)

### <u>Glycogen</u>

- 1 a polysaccharide;
- 2 (chain of)  $\alpha$ /alpha glucose;
- 3 <u>1-4</u> with (some) <u>1-6 links</u>/branches;
- 4 insoluble;
- 5 (energy) store;
- 6 stored/produced in muscle/liver;

## <u>Glucagon</u>

- 1 a globular protein;
- 2 (chain of) amino acids;
- 3 ref. to peptide bonds;
- 4 soluble in plasma;
- 5 produced by  $\alpha$ /alpha cells in pancreas/islets of Langerhans;
- 6 a hormone/regulates/raises blood glucose;

- 1 ref. energy reserves;
- 2 ref. lipids buoyancy;
- 3 ref. lipids (thermal) insulation;
- 4 ref. lipids more energy per unit <u>mass</u>/higher calorific value;
- 5 ref. to metabolic water;
- 6 ref. protection;

(ref) a continuous piece/sequence

### **QUESTION 9**

#### (a)

- 1 length of DNA;
- 2 gene/ Lacl, coding for repressor/inhibitor protein;
- 3 ref. to CAP (catabolic activator protein) binding site next to promoter;
- 4 promoter (gene);
- 5 RNA polymerase binding site;
- 6 ref. to operator;
- 7 binding site for repressor/inhibitor protein;
- 8 ref. to (3) <u>structural</u> gene;
- 9 Lac Z (galactosidase);
- 10 Lac Y (lactose permease);
- 11 Lac A (lactose transacetylase);
- 12 ref. to termination sequence;

8 max

## (b)

- 1 lactose enters (bacterial) cell; basal level of lac permease
- 2 ref. to method of entry of lactose / qualified leaky membrane / small amount in;
- 3 allolactose / lactose binds to repressor protein;
- 4 results in a change in shape of repressor protein;
- 5 repressor protein no longer attached (to DNA);
- 6 exposes operator site;
- 7 allows RNA polymerase activity;
- 8 transcription (of genes coding for proteins required to utilise lactose);
- 9 leading to synthesis of proteins;
- 10 ref. named enzymes/proteins;

- gene switching / majority of genes switched off / for regulation of gene expression;
- 2 so that bacterium only produces enzymes required;
- 3 <u>inefficient</u> / ref. waste of energy/resources **A converse**
- 4 ref. to selective advantage / disadvantage;
- 5 ref. group genes together for control;
- 6 ref. to bacteria able to use a <u>variety</u> of sugars/substrates

# Paper 3 (Applications)

### **QUESTION 1**

(a)

(i)

bacterium/bacteria;

### ignore correct qualification (e.g.named)

### (ii)

break phosphodiester bonds within polynucleotide strands/DNA; (R) cut at, restriction site/gene

## (iii)

protection;

breaks down, foreign/viral/phage, DNA;

allow ecf from (i) 1 max

(b)

- (i)
- 1 (BamHI + HindIII + EcoR1) all produce, sticky ends/cohensive ends/overhangs/staggered ends;
- 2 allows, cut DNA/gene, to be, joined/annealed/spliced, to another DNA; ignore vector
- 3 using complementary basepairing;
- 4 Alul + Haelll require additional procedure to add on, sticky ends/linker DNA/caps; ora 3 max

(ii)

- 5' represents end (of strand with nucleotide) with 'free' carbon 5 (on deoxyribose);
- 2 3' represents end (of strand with nucleotide) with 'free' carbon 3 (on deoxyribose);
- 3 5' has phosphate/ 3' has hydroxyl;
- 4 DNA strands are antiparallel;

(c)

(i)

# genomic

- 1 DNA is broken (into fragments) using Restriction Enzyme(s); Reject: treated
- 2 fagments/DNA, incorporated into vector;
- 3 X phage/plasmid/cosmid/BAC/YAC; (BAC: bacteria artificial chromosome, YAC: yeast artificial chromosome)
- 4 create, further copies/clones, of DNA;

## cDNA

- 5 start with mRNA;
- 6 to DNA using Reverse Transcriptase;
- 7 make double-stranded using DNA polymerase;

4max for any section

### (ii)

the cDNA Library

- 1 smaller/ more compact;
- 2 contains only, coding sections (of DNA)/ exons;
- 3 <u>introns</u> have been removed/ no junk <u>DNA;</u>
- 4 easier to find desired gene (DNA clone)/no need for gene probes;
- 5 contains genes that are being expressed/reflects proteins actually present (in cell);
- 6 genes are intact;
  - (R) not been cleaved

## credit reverse statements for genomic library 3 max

#### (a)

### Stage 1

- denatures / H bonds broken: 1
- (2) DNA strand separate; 2
- A:  $90 100^{\circ}C$ 3 by heating to  $95^{\circ}C$ ; \*

## Stage 2

- primer / oligonucleotides, bind / join to single (DNA) strands / 3' ends; 4
- 5 annealing:
- by cooling to 54<sup>0</sup>C: \* A:  $30 - 65^{\circ}C$ 6

## Stage 3

- new strands (of DNA) are synthesized by, TAQ / DNA polymerase; 7
- starts at positions of primers / oligonucleotides; 8
- A:  $60 75^{\circ}C$ by heating to  $72^{\circ}C$ ; 9

# (b)

**(i)** 2<sup>26</sup> / 67108864

Look for 2<sup>26</sup> first

# (ii)

- (may only be) very small amount (of original DNA) to work with; 1
- can, amplify / produce many copies / produce very large amounts of DNA / 2 genetic material;
- 3 (large amounts) make analysis easier:

# (C)

- DNA sample / genome (removed from organism); 1
- 2 RFLP / Restriction Fragment Length Polymorphism;
- 3 (sample DNA) cut with restriction enzymes,
- 4 gel electrophoresis / described;
- radioactive probe + X-ray film or ethidium bromide / stain + UV light (to locate 5 bands position)
- wild and captive populations have different genetic profiles; 6
- 7 genetic fingerprints of captive individuals obtained;
- 8 compare fingerprint / DNA, with that of organism being investigated;
- 9 if bred in captivity there will be a match in, banding / pattern; ora

## Comments by examiner:

- Students were weak in identifying how genetic fingerprinting is used to verify what type of organism (wild / bred in captivity) as many elaborated too much on the
- Genetic Fingerprinting method.
- Restriction Enzymes are not just added to the mixture but also used to 'cut' DNA

#### (a)

(i)

- derived from one parent/not a product of sexual reproduction ; (ignore asexual reproduction)
- 2 products of mitotic division;
- 3 genetically identical/share same DNA/same genotype/same genome/ same alleles; (R) same genes/similar

## (ii)

- 1 much larger numbers can be produced/mass production ;
- 2 requires less space;
- 3 (cultures) easy to transport/ref. air freight;
- 4 ensures new plants are disease-free;
- 5 genetic modifications possible;
- 6 not limited by seasons/external conditions/weather; (R) faster techniques

# (b)

<u>2,4-D</u>

- 1 optimum peak/highest/described, is 0.3mg dm<sup>-3</sup> with growth rate of 73/74;
- 2 idea that slight change from optimum is critical;

## <u>Cytokinin</u>

- $\overline{3}$  optimum peak/highest/described, is 8(.0)mg dm<sup>-3</sup> with growth rate of 50;
- 4 not critical/large change from optimum to results in smallest decrease in growth (rate);

<u>Thiamine</u>

- 5 optimum peak/highest/described, is 1(.0)mgdm<sup>-3</sup> with growth rate of 61/62;
- 6 marked effect/critical from 0.8mg mg dm $^{-3}$ ;
- 7 2,4-D has highest growth rate/cytokinin has lowest growth rate;
- 8 cytokinin required at highest concentration / 2,4-D required at lowest concentration;
- 9 ref. qualified pattern of variation for one component;
- 10 use a mixture of all 3 at optimal levels;
- 11 idea of possible interaction between components;

## Comments by examiner:

- Arbitrary units not required in this question but many not be true for all questions.
- Guide students to state the complete units in all questions wherever possible.
- Need to observe scale of graph
- Some form of comparison between graphs wanted by examiner though not very clearly stated in question

#### (c) (i)

- 1 unable to fix nitrogen;
- 2 to produce, amino acids/proteins;
- 3 e.g. of required protein ; (e.g. enzymes/cytochrome/etc)
- 4 DNA/RNA/nucleic acid/nitrogenous base/named base/nucleotide;
- 5 ATP;
- 6 chlorophyll;

(ii)

- ammonium nitrate contains more nitrogen;
- 2 may be, easier to assimilate/easier to absorb/more accessible, ammonium (ion rather than potassium ion);

#### (a)

<u> SĆID</u>

- 1 more males affected (than females)
- 2 recessive
- 3 (most common form is) sex-linked (X-chromosome)
- 4 mutation of IL2RG (interleukin-2 receptor gamma) gene (for gamma chain on lymphocyte receptor)
- 5 ADA (adenosine deaminase) deficiency
- 6 mutation of gene on chromosome 20
- 7 prevents formation of , lymphocytes / T cells / B cells
- 8 toxic metabolites not broken down so lymphocytes killed
- 9 so susceptible to opportunistic infections (award only if S7 and S8 awarded)
- 10 children usually die (if not treated)
- 11 pneumonia / PCP / meningitis / chicken pox / measles / other valid

### Cystic fibrosis

- 1 autosomal / chromosome 7
- 2 recessive
- 3 mutation of, CFTR / CFTCR gene
- 4 deletion of, 3 base (pairs) / codon
- 5 CFTR / CRTCR, protein has lost phenylalanine
- 6 (channels) no longer able to transport chloride ions
- 7 causing thick mucus
- 8 resulting in stated problem in, lungs / pancreas / reproductive system
- 9 reduces life expectancy

### (b)

- 1 gene therapy is a technique for introducing 'normal', allele/gene; *if "therapeutic"* need to give further explanation
- 2 new, allele/gene, generates, functional/correct, protein product; accept specific example of protein product
- 3 restores target cell to normal state / corrects function of cell / corrects phenotype;

## Viral delivery systems

- 1 viruses are genetically engineered to carry 'normal' human gene;
- 2 adenoviruses/retroviruses/MMLV (for SCID);
- 3 virus releases genetic material into target cells;
- 4 in SCID the gene is inserted (by viruses) into, bone marrow cells / blood stem cells / lymphocytes, *ex vivo* / *in vitro*;
- 5 in cystic fibrosis the gene is inserted into cells within body;

#### Non-viral delivery systems

- 1 direct introduction of DNA into target cells (gene gun or other direct method e.g. micropipette);
- 2 creation of, lipid sphere / liposome, which carries DNA;
- 3 liposome can / use of electroporation to, pass through target cell membrane; must be linked to pt 10
- 4 link DNA to molecule that will bind to receptors in cell membrane;
- 5 DNA is engulfed by membrane;
- 6 possibility of introducing 47<sup>th</sup> chromosome into cells;

- difficult to get DNA to integrate into target cell genome;
- 2 results can be short-lived / patients need to be treated on a frequent basis;
- 3 problem with controlling the activity of gene expression;
- 4 risk of stimulating immune / allergic response;
- 5 virus vector may, regain / develop, virulence;
- 6 incorrect insertion of gene may cause, named / described, problem to patient;
- 7 possible problem with finding vector for large gene;
- 8 many genetic diseases are a result of presence of many defective genes / multiallelic / multigene conditions;
- 9 impossible to introduce many genes at the same time;
- 10 problem dealing with, dominant condition / non-dividing cells;
- 11 ref. problem with liposomes; (toxicity / less efficiency);