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2214-6007 - 6 - M14/4/BIOLO/HPM/ENG/TZ1/XX 14. The diagram shows a simplifi ed Lake Michigan foodweb. Which pair of terms describes the trophic

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MARKSCHEME
2214-6007 - 5 - Turn over M14/4/BIOLO/HPM/ENG/TZ1/XX 10. What is the result after the irst meiotic division of one diploid animal cell containing

BIOLOGY HIGHER LEVEL PAPER 1
M14/4/BIOLO/HPM/ENG/TZ2/XX 10. A pea plant that is homozygous for purple lowers is crossed with a pea plant having white lowers and the descendants are all plants with purple lowers. One of these F1 plants is then crossed with a pea plant having white lowers. In the resulting offspring, what can be expected? A. 100 % of plants with purple lowers

Friday 9 May 2014 (afternoon)
- 8 - M14/4/BIL/HPM/ENG/T2/XX 13. How can forensic scientists obtain sufficientdata from one hair follicle to make a reliable identification by DNA profiling? A. By performing PCR with DNA from the sample B. By digesting the sample with more than one restriction enzyme C. By performing electrophoresis with many other known samples D.

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Markscheme - Weebly
motor end plates - shown as widened ends of multiple branches of the nerve fibre; [4 max] Reject the points for axon and myelin sheath if the labelling line does not clearly point to the structure.

MARKSCHEME
- 3 - Turn over CAMG0 4. What structure is labelled X? A. DNA B. Beta pleated sheet C. Alpha helix D. Prosthetic group 5. In the experiments performed by Meselson and Stahl, were grown for many generations E. coli in 15N then for one generation in 14N.What results for the DNA of the last generation showed that

Biology Higher and standard level
C. 4 4 D. 4 2 13. The allele for tall T is dominant to the allele for dwarf t. Which of the following represents a test cross? A. x tttt B. TT x Tt C. x tTt D. x TtT 14. Which process can be used to amplify small fragments of DNA? A. Gel electrophoresis B. Polymerase chain reaction C. DNA profiling D. Electron microscopy M15/4/BIOLO/PM/ ENG ...

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(c) a. complete human DNA/chromosomes sequenced; b. identification of all human genes / find position/map (all) human genes ; c. find/discover protein structures/functions; d. find evidence for evolutionary relationships/human origins/ancestors; e. find mutations/base substitutions/single nucleotide polymorphisms; f. find genes causing ...

Full text of "Biology_HL"
opped to incorrec (see point 4). • Where M and A marks are noted on the same line, eg M1A1 , this usually means M1 for an attempt to use an appropriate method (eg substitution into a formula) and A1 for using the

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