



ANSWER BOOK

Genomes, screening and gene therapy: what a sequence is good for

Gene regulation, genomics and biotechnology · Unit B17

AQA 3.8.3, 3.8.4.2, 3.8.4.3 · OCR A 6.1.3 · Edexcel A Topic 2, Topic 6 · CAIE 19.2

5 questions · 16 marks

NAVY EDITION

SECTION A · Short, 1 to 2 marks

- A1** **B1** a sequence of bases for the organism, coding and non-coding alike. **B1** [2]
annotation locates the genes within that sequence and works out which non-coding regions do something, and it is a separate and slower job done afterwards.
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SECTION B · Standard, 3 to 5 marks

- B1** **B1** bacterial genes are continuous, with no introns, so one gene gives one [4]
polypeptide and the coding sequence can be read straight off the DNA. **B1** human
genes contain introns, which have to be identified and removed before the coding
sequence is known. **B1** alternative splicing lets one human gene give several different
polypeptides. **B1** large stretches of the human genome are regulatory rather than
coding, and post-translational modification changes the polypeptide again after
translation.
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- B2** **B1** somatic therapy alters body cells such as bone marrow, retina or airway [4]
epithelium, whereas germ line therapy alters a gamete, a zygote or an early embryo. **B1**
a somatic change is not inherited, because the cells that become sperm or eggs are
untouched, whereas a germ line change is present in every cell of the resulting person,
including those cells, and passes to every descendant. **B1** somatic treatment often has
to be repeated, because cells that are not stem cells are replaced within weeks,
whereas a germ line change is permanent and cannot be withdrawn from the family
line. **B1** somatic therapy is licensed for several conditions, whereas germ line therapy in
a pregnancy is prohibited by law in every country that has legislated on the question.
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- B3** **B1** the probe is a short single strand of DNA whose base sequence is [3]
complementary to the sequence being looked for, carrying a radioactive or a
fluorescent label. **B1** the sample DNA is made single-stranded by heating and the
probe is added, and hybridisation occurs wherever the probe meets a complementary
sequence, held by hydrogen bonds between complementary bases. **B1** unbound probe
is washed away, and the label is then detected, on photographic film or under
ultraviolet light, only where the probe has bound.
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- B4** **B1** the quoted figure is the probability of obtaining a matching profile given [3]
that the person is not the source of the sample. **B1** the lawyer has reversed it into the
probability that the person is not the source given that the profile matches, which is a
different quantity and is not what the figure measures; reversing the two is the
prosecutor's fallacy. **B1** the figure also assumes an unrelated person, and relatives
share far more alleles than strangers, while a degraded sample gives a partial profile
at fewer loci and the match probability weakens by orders of magnitude.
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