

3.5 Transcription and translation – *summary of mark schemes*

3.5.1	<p>Compare the structure of RNA and DNA.</p> <p>Mark Scheme</p> <ul style="list-style-type: none">A. deoxyribose versus ribose;B. thymine versus uracil;C. two strands versus one / double helix versus single strand;
3.5.2	<p>Outline DNA transcription in terms of the formation of an RNA strand complementary to the DNA strand by RNA polymerase.</p> <p>Mark Scheme</p> <ul style="list-style-type: none">A. RNA polymerase controls transcription / is the enzyme used in transcription;B. DNA is unwound by RNA polymerase;C. DNA is split into two strands;D. mRNA is made by transcription;E. promoter region (by start of gene) causes RNA polymerase to bind;F. anti-sense / template strand of DNA is transcribed;G. direction of transcription is 5' → 3';H. free nucleotide triphosphates used;I. complementary base pairing between template strand and RNA nucleotides / bases;J. RNA contains uracil instead of thymine;K. terminator (sequence) stops RNA polymerase / transcription;L. mRNA is released / RNA polymerase released;
3.5.3	<p>Describe the genetic code in terms of codons composed of triplets of bases.</p> <p>Mark Scheme</p> <ul style="list-style-type: none">A. composed of mRNA base triplets;B. called codons;C. 64 different codons;D. each codes for the addition of an amino acid to a growing polypeptide chain;E. the genetic code is degenerate;F. meaning more than one codon can code for a particular amino acid;G. the genetic code is universal;H. meaning it is the same in almost all organisms;I. (AUG is the) start codon;J. some (nonsense) codons code for the end of translation;
3.5.4	<p>Explain the process of translation, leading to polypeptide formation.</p> <p>Mark Scheme</p> <ul style="list-style-type: none">A. consists of initiation, elongation and termination;B. mRNA is used as a template / guide;C. mRNA translated in a 5' to 3' direction;D. binding of ribosome to mRNA;E. small sub-unit then large;F. first / initiator tRNA binds to start codon / to small subunit of ribosome;G. AUG is the start codon;H. second tRNA binds to ribosome;I. large subunit moves down mRNA after a second tRNA binds;J. mRNA "read" in base triplets / codon;K. each codon specifying addition of a particular amino acid to the growing polypeptide;L. tRNA bring amino acids (to mRNA-ribosome complex);M. amino acid / polypeptide on first tRNA is transferred / bonded to amino acid on second tRNA;N. peptide bonds between amino acids / peptidyl transferase;O. requires GTP;P. movement of ribosome / small subunit of ribosome down the mRNA;Q. loss of tRNA and new tRNA binds;R. reach a stop codon / termination;

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| | <ul style="list-style-type: none">S. stop codon has no corresponding tRNA / amino acid / causes release of polypeptide;T. polypeptide released;U. tRNA activating enzymes link correct amino acid to each tRNA;V. (activated) tRNA has an anticodon and the corresponding amino acid attached;W. many ribosome / polyribosomes bind to same mRNA; |
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