系所班組別:生命科學院甲組、醫學生物科技學程 考試科目(代碼):分子生物學(0404、0704)

共__9__頁,第__1__ *請在【答案卷】作答

I 單選題 (每題 2 分,共 56 分)

- 1. Biochemical and genetic experiments have demonstrated that the ______ of tRNA are important for recognition by its cognate aminotransferase-tRNA synthetase.
 - A. acceptor stem and anticodon loop
 - B. T loop and variable loop
 - C. anticodon loop and T loop
 - D. variable loop and D loop
 - E. D loop and T loop
- 2. Which of the following molecules does not resemble a tRNA molecule?
 - A. tmRNA
 - B. EF-Tu
 - C. puromycin
 - D. RRF
 - E. suppressor tRNA
- 3. Which of the following antibiotics does not inhibit protein synthesis by binding to the ribosome?
 - A. chloroamphenicol
 - B. streptomycin
 - C. ampicillin
 - D. erythromycin
 - E. tetracycline

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共__9__頁,第__2__頁 *請在【答案卷】作答

- 4. Put the following steps of prokaryotic translation initiation in the correct order.
 - (1) Binding of IF1, IF2, and GTP to the 30S subunit.
 - (2) Binding of IF3 to the 30S subunit.
 - (3) Binding of the 50S subunit and loss of IF1 and IF3.
 - (4) Dissociation of the 70S ribosome.
 - (5) Formation of the 70S initiation complex by dissociation of IF2 and GTP hydrolysis.
 - (6) Formation of the 30S initiation complex.
 - A. 1,2,3,4,5,6
 - B. 2,1,6,3,5,4
 - C. 5,6,3,2,1,4
 - D. 4,2,1,6,3,5
 - E. 2,4,1,6,3,5
- 5. Please place the steps of translation elongation in the correct order.
 - (1) Peptidyl transferase forms a peptide bond between the peptide in the P site and the newly arrived aminoacyl-tRNA in the A site.
 - (2) EF-G, with GTP, translocates peptidyl-tRNA to the P site.
 - (3) EF-Tu, with GTP, binds an aminoacyl-tRNA to the ribosomal A site.
 - A. 3, 2, 1
 - B. 1, 2, 3
 - C. 2, 3, 1
 - D. 1, 3, 2
 - E. 3, 1, 2
- 6. Which of the following evidence can be considered as supports or necessity for discontinous synthesis of one strand of double-stranded DNA molecules?
 - A. the presence of Okazaki fragments as replication intermediates
 - B. DNA polymerase requires a primer
 - C. DNA replicates unidirectionally
 - D. DNA replicates bidirectionally

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共__9__頁,第__3__ *請在【答案卷】作答

- 7. Which one concerning hybrid dysgenesis in *Drosophila* is true?
 - A. Hybrid dysgenesis occurs when the P element is contributed by the egg and sperm is devoid of P element.
 - B. Hybrid dysgenesis occurs when both parents (egg and sperm donors) carry P element.
 - C. Transposition of P element is responsible for killing somatic cells of F1 hybrid
 - D. The molecular basis of hybrid dysgenesis lies at the tissue-specific splicing of the primary transcript of transposase
- 8. Which one of the following process requires RecA activity
 - A. activating immunoglobulin gene rearrangement
 - B. resolving a Holliday junction during recombination
 - C. activating SOS gene expression
 - D. Integration of λ phage genome to bacterial chromosome
- 9. Which of the following enzyme uses RNA to synthesize DNA?
 - A. Klenow enzyme
 - B. RNA replicase
 - C. telomerase
 - D. Escherichia coli DNA polymerase IV
- 10. Which statement is <u>not true</u> for class I and III promoters in the eukaryotes?
 - A. For eukaryotic RNA polymerase III-dependent promoters (class III promoter), the TATA box is recognized by the general transcription factor TFIIIB.
 - B. For class I promoter in human, the core-binding factor SL1 is required to recruit RNA polymerase I.
 - C. TFIIIB acts as an assembly factor that binds to the internal promoter and helps TFIIIA to bind to a region just upstream of the transcription start site.
 - D. TFIIIA is required for transcription of the 5S rRNA genes, but not for the tRNA genes.
 - E. Although class I promoters are variable in sequence across species, they do contain the AT-rich initiator with a conserved sequence.

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共__9__頁,第__4__頁 *請在【答案卷】作答

- 11. Please choose one <u>wrong</u> statement for the roles of different general transcription factors in eukaryotic RNA polymerases.
 - A. TATA-binding factor (TBP) is also required for transcription in the Archaea.
 - B. The TBP-associated factor 1 (TAF1) serves as an assembly factor around which other TAFs can aggregate and also contains a histone acetyltransferase activity.
 - C. TFIIB is a general transcription factor with its C-terminal domain to interact with RNA polymerase II and can determine the direction of transcription.
 - D. TBP-free TAF-containing complex (TFTC) can sponsor the formation of preinitiation complex of RNA polymerase II without the presence of TFIID.
 - E. The negative cofactor 2 (NC2) can stimulate RNA polymerase II-related gene transcription from downstream promoter element (DPE)-containing promoters.
- 12. Which of the following statement is true for gene transcription in the eukaryotes?
 - A. Tfg2 is the second largest subunit of TFIIF and is homologous to bacterial alpha-subunit of RNA polymerase.
 - B. Mediator is a collection of proteins that is a part of most RNA polymerase II preinitiation complexes and is not required for transcription initiation, but is required for activated transcription.
 - C. TFIIIA is a general transcription factor for RNA polymerase III and contains 9 zinc finger structures in which one finger containing 6 cycteines to coordinate 2 zinc ions.
 - D. In the absence of ligand, the thyroid hormone receptor exists as inactive form in cytoplasm complexed with heat shock protein 90.
 - E. Silencers are DNA-binding proteins that can cause strong bending in DNA.
- 13. Please choose one of the following technique that <u>cannot</u> be used to quantify the level of gene expression.
 - A. Nuclear run-off assay.
 - B. Northern blot analysis.
 - C. S1 mapping.
 - D. RFLP analysis.
 - E. RNase protection.

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共_9_頁,第_5_頁 *請在【答案卷】作答

- 14. Choose one <u>wrong</u> statement from the followings regarding eukaryotic transcription?
 - A. In flowing plants, RNA polymerase IV and V are involved in gene silencing, which are performed by RNA polymerase II in other eukaryotes.
 - B. TFIIA can interfere with the interaction between TAF1 of RNA polymerase II N-terminal domain & DNA-binding surface of TBP, freeing up TATA-binding protein (TBP) for promoter binding.
 - C. Rpb10 and Rpb12 are common subunits in all three yeast polymerases.
 - D. An enhanceosome is a collection of DNA sequences (enhancers) that can enhance transcription by themselves.
 - E. The incoming nucleotide first encounters the E ("entry") site and the "A" site of RNA polymerase II is where a phosphodiester bond forms.
- 15. A DNA element found in prokaryotes that binds tightly to a specific repressor and thereby regulates the expression of adjoining genes is called
 - A. Enhancer
 - B. Promoter
 - C. Terminator
 - D. Suppressor
 - E. Operator
- 16. Which of the following is a possible mechanism for the action of NusA in termination?
 - A. It binds to NusB to promote detachment of the RNA polymerase for the DNA template.
 - B. It interacts with S10 to promote dissociation of the RNA polymerase.
 - C. It binds with N to promote hairpin loop formation.
 - D. It stimulates termination at intrinsic terminator by facilitating hairpin loop formation.
 - E. It binds to the RNA polymerase causing it to stall.

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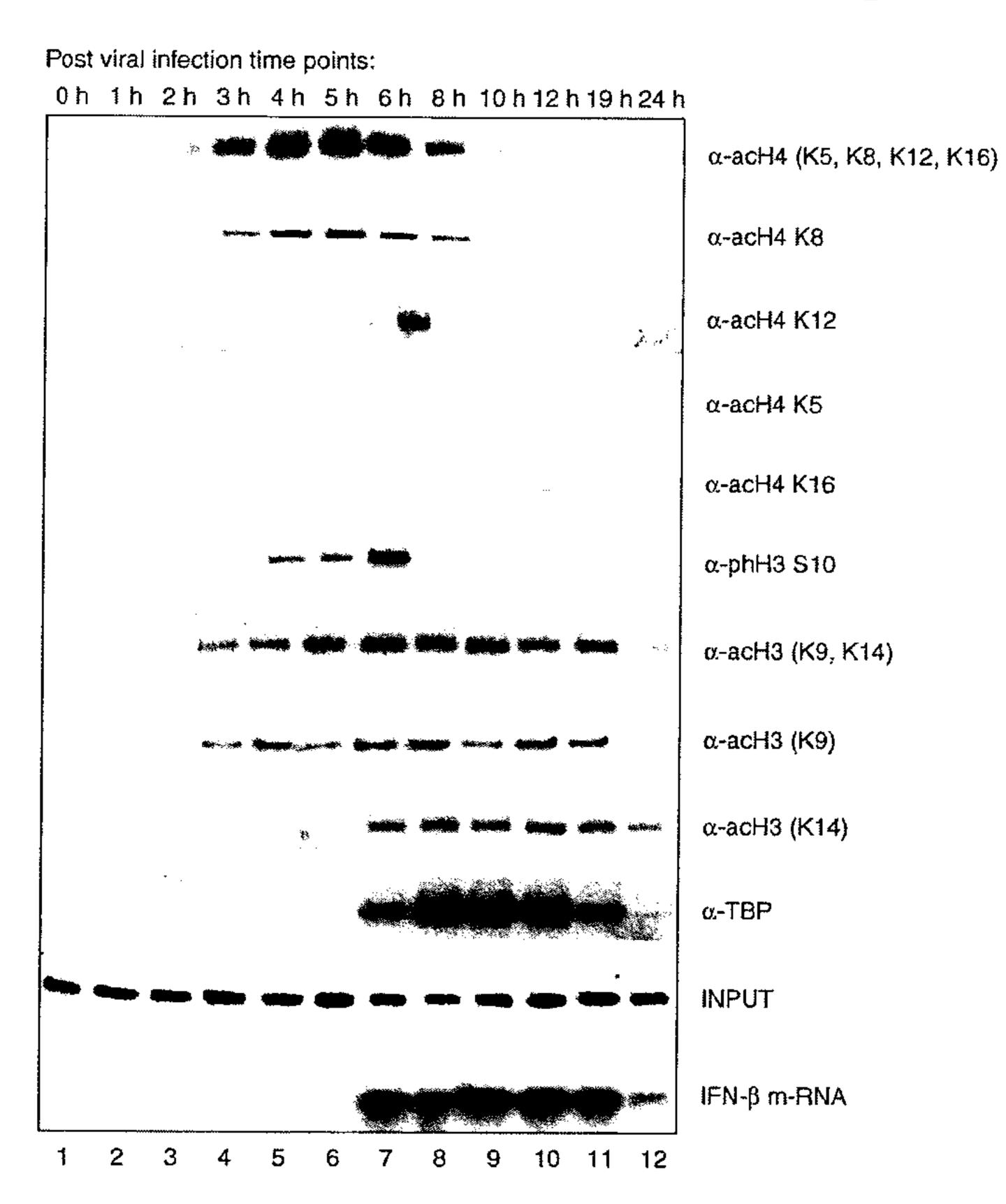
共__9__頁,第__6__ *請在【答案卷】作答

- 17. According to the result of affinity labeling, which one of the following subunits lies near the active site of bacterial RNA polymerase where phosphodiester bonds are formed?
 - A. α
 - Β. β
 - C. β'
 - D. δ
 - E. ω
- 18. Which of the following statements is not correct about the cro gene?
 - A. It is adjacent to the *cIII* gene.
 - B. It must be repressed during lysogeny.
 - C. It belongs to the immediate early gene.
 - D. It must be stimulated during the lytic cycle.
 - E. Its product represses repressor (CI) activity.
- 19. Which of the following techniques is most useful in determining whether the α -subunit of E. coli RNA polymerase is involved in contacting a promoter UP element?
 - A. Run-off transcription assay.
 - B. Southern analysis.
 - C. DNase footprinting experiment.
 - D. Filter binding assay.
 - E. DNA sequencing.
- 20. Which of the following explains the events in late stage of phage SPO1 infection in *B. subtilis*?
 - A. There is a high level of transcription of host genes.
 - B. Host polymerase holoenzyme directs transcription.
 - C. Phage SPO1-specific core polymerase participates in transcription.
 - D. The host core enzyme participates in transcription in conjunction with the phage-encoded specific sigma factor.
 - E. The host sigma factor specifies transcription of genes.

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共__9__頁,第__7__頁 *請在【答案卷】作答

- 21. Which chromatin folding is <u>unlikely</u> to occur in the putative 30-nm chromatin fiber?
 - A. solenoid model
 - B. stem-loop model
 - C. Zig-Zag model
 - D. two-start double helical model
- 22. What is **NOT** a proposed role of Histone H1?
 - A. nucleosome stabilizer
 - B. inhibitor of transcription
 - C. a linker histone
 - D. a component of the core nucleosome
- 23. Chromatin Immunoprecipitation (ChIP) analysis shown below revealed the timing of histone acetylation in chromatin at INF-β promoter of HeLa cells after Sendai virus infection. Which hypothesis may **NOT** be true?



- A. Phosphorylation of histone H3 S10 is necessary for lysine 14 acetylation.
- B. Acetylation of lysine 8 of histone H4 promotes SWI/SNF complex binding to nucleosome.
- C. Acetylation of lysine 14 of H3 is required to recruit TBP to INF-beta promoter.
- D. Acetylation of lysine 16 of H4 is required for phosphorylation on S10 of histone H3.

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共__9__頁,第__8__ *請在【答案卷】作答

- 24. What is the consensus sequence of mammalian pre-mRNA intron?
 - A. exon/GU-intron-AG/exon
 - B. exon/UU-intron-GG/exon
 - C. exon/CA-intron-GG/exon
 - D. exon/GA-intron-AA/exon
- 25. Which of the following enzymes is **NOT** involved in the process of vaccinia virus and reovirus mRNA cap synthesis?
 - A. Methyltransferase
 - B. RNA triphosphatase
 - C. Guanylyl transferase
 - D. RNA dependent protein kinase
- 26. Which eukaryotic molecule contains a poly(A) tail during their biogenesis?
 - A. 5S rRNA
 - B. tRNA
 - C. U6 snRNA
 - D. mRNA
- 27. Which of the following descriptions about polyadenylation signal in mammalian cells is true?
 - A. AAUAAA is the most frequent and active sequence.
 - B. Polyadenylation signal is followed 23-24 bp later by a UC-rich motif and then by a C-rich motif.
 - C. The cleavage/ polyadenylation site is within the polyadenylation signal sequence.
 - D. The cleavage/polyadenylation site is within the U-rich motif.
- 28. Which of the following RNA processing events does **NOT** take place while eukaryotic gene transcription is underway?
 - A. splicing
 - B. capping
 - C. polyadenylation
 - D. cytoplasmic transportation

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共_9_頁,第_9_頁 *請在【答案卷】作答

II 填充題 (每題 2 分,共 4 分)

1. _____ repair system is responsible for maintaining the stability of microsatellite. Failure of this repair may increase the chance of cancer in human.

2. Many of the DNA polymerases contain _____ activity which is responsible for maintaining the accuracy of DNA replication.

III 問答題 (共 40 分)

- 1. What would be the effect on a G protein's activity of (a) inhibiting its GAP? (b) inhibiting its guanine nucleotide exchange protein? Give your explanation. (5 points)
- 2. What is inducible expression vector? What benefit of using it in producing recombinant proteins? (5 points)
- 3. Consider two theoretical eukaryotic transposons, A and B. Each contains an intron. Each transposes to a new location in the yeast genome and then is examined for the presence of the intron. In the new locations, you find that A has no intron, but B does. What can you conclude about the mechanisms of transposition movement for A and B from these facts? (3 points)
- 4. How the immune systems of vertebrates can produce many millions of different antibodies? (5 points)
- 5. The reverse transcription (RT) followed by real-time quantitative PCR is the technique of choice to analyze mRNA expression. (a) Please <u>describe</u> the <u>principle</u> for the real-time quantitative PCR (5 points) (b) Please use the 2^{-\times_Ct} method to <u>explain</u> how real-time quantitative PCR can be used to obtain <u>a relative expression</u> level of the same gene from two different biosamples. (5 points)
- 6. (a) Present a model to explain attenuation in the *trp* operon in *E. coli*. (3 points) (b) How is *trp* attenuation overridden in *E. coli* when tryptophan is scarce (缺乏的)? (5 points)
- 7. Describe Drosha and Mirtron pathways for human miRNA biogenesis. (4 points)